GenCore version 4.5

OM nucleic - nucleic search, using sw model

Run on: August 28, 2002, 08:01:00 ; Search time 2751.06 Seconds
(without alignments)
12197.381 Million cell updates/sec

Title: US-08-711-417C-165
Perfect score: 1551
Sequence: 1 ATGGATGCTGACGAGGGTCA.....ACCGCTTCCACATGAGCTAA 1551

Scoring table: IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Searched: 21979536 seqs, 10817449327 residues

Total number of hits satisfying chosen parameters: 43959072

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	Length	DB	ID	Description
1	1551	100.0	1788	18	US-09-435-327A-19	Sequence 19, Appl
2	1497.8	96.6	6171	56	US-60-172-373-18124	Sequence 18124, A
3	1468.8	94.7	3629	14	US-09-023-655-923	Sequence 923, App
4	1468.8	94.7	3629	18	US-09-435-327A-18	Sequence 18, Appl
5	1468.8	94.7	3629	18	US-09-442-384A-721	Sequence 721, App
6	1468.8	94.7	4786	25	US-09-649-161-10565	Sequence 10565, A
7	1468.8	94.7	4786	25	US-09-652-816-9158	Sequence 9158, Ap
8	1468.8	94.7	4786	27	US-09-698-013-6274	Sequence 6274, Ap
9	1468.8	94.7	4786	27	US-09-698-014-4890	Sequence 4890, Ap
10	1468.8	94.7	4786	28	US-09-717-350-5922	Sequence 5922, Ap
11	1468.8	94.7	4786	29	US-09-726-802-2123	Sequence 2123, Ap
12	1468.8	94.7	4786	29	US-09-726-811-4096	Sequence 4096, Ap
13	1457.8	94.0	6203	71	US-60-324-185-24401	Sequence 24401, A
14	1386	89.4	1386	1	PCT-US99-02559-16	Sequence 16, Appl
15	1386	89.4	1386	6	US-08-238-212A-3	Sequence 3, Appli
16	1386	89.4	1386	8	US-08-465-590B-3	Sequence 3, Appli
17	1386	89.4	1386	11	US-08-733-622A-16	Sequence 16, Appl
18	1386	89.4	1386	11	US-08-733-622B-16	Sequence 16, Appl
19	1386	89.4	1386	14	US-09-019-348-16	Sequence 16, Appl
20	1386	89.4	1386	14	US-09-019-348A-16	Sequence 16, Appl
21	1386	89.4	1386	29	US-09-755-830-2	Sequence 2, Appli
22	1382.4	89.1	1611	5	US-08-121-438-3	Sequence 3, Appli
23	1174.8	75.7	2049	1	PCT-US99-02559-18	Sequence 18, Appl
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27	1174.8	75.7	2049	11	US-08-733-622B-18	Sequence 18, Appl
28	1174.8	75.7	2049	14	US-09-019-348-18	Sequence 18, Appl
29	1174.8	75.7	2049	14	US-09-019-348A-18	Sequence 18, Appl
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39	837	54.0	1542	75	US-60-360-207-1196	Sequence 1196, Ap
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44	813.2	52.4	1788	8	US-08-465-590B-2	Sequence 2, Appli
45	813.2	52.4	1788	11	US-08-733-622A-15	Sequence 15, Appl

ALIGNMENTS

RESULT 1
US-09-435-327A-19
; Sequence 19, Application US/09435327A
; GENERAL INFORMATION:
; APPLICANT: Uckun, Fatih M.
; APPLICANT: Crotty, Mya L.
; TITLE OF INVENTION: IKAROS ISOFORMS AND MUTANTS
; FILE REFERENCE: 12152.35USU1
; CURRENT APPLICATION NUMBER: US/09/435,327A
; CURRENT FILING DATE: 1999-11-05
; PRIOR APPLICATION NUMBER: 60/107,229
; PRIOR FILING DATE: 1998-11-05
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 19
; LENGTH: 1788
; TYPE: DNA
; ORGANISM: Homo sapiens
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Query Match 100.0%; Score 1551; DB 18; Length 1788;
Best Local Similarity 100.0%; Pred. No. 0;
Matches 1551; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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RESULT   2
US-60-172-373-18124
; Sequence 18124, Application US/60172373
; GENERAL INFORMATION:
;  APPLICANT: Morris, MacDonald
;  APPLICANT: Lal, Preeti
;  APPLICANT: Diep, Dinh
;  TITLE OF INVENTION: Method for the Identification of Sequence Polymorphisms Using
;  TITLE OF INVENTION: Polynucleotide Sequence Databases, and Single Nucleotide Polymorp
;  FILE REFERENCE: GX-0006 P
;  CURRENT APPLICATION NUMBER: US/60/172,373
;  CURRENT FILING DATE: 1999-12-16
;  NUMBER OF SEQ ID NOS: 25,772
;  SOFTWARE: PERL Program
; SEQ ID NO 18124
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;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: misc_feature
;   OTHER INFORMATION: Incyte ID No: 345320.3
;   FEATURE:
;   NAME/KEY: unsure
;   LOCATION: 3401-3583
;   OTHER INFORMATION: a, t, c, g, or other
US-60-172-373-18124


  Query Match             96.6%;  Score 1497.8;  DB 56;  Length 6171;
  Best Local Similarity   98.5%;  Pred. No. 1.6e-303;
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Db    562 ggagaacggcccttccagtgcaatcagtgcggggcctcattcacccagaagggcaacctg 621

Qy    481 CTCCGGCACATCAAGCTGCATTCCGGGGAGAAGCCCTTCAAATGCCACCTCTGCAACTAC 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    622 ctccggcacatcaagctgcattccggggagaagcccttcaaatgccacctctgcaactac 681

Qy    541 GCCTGCCGCCGGAGGGACGCCCTCACTGGCCACCTGAGGACGCACTCCGTTGGTAAACCT 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||

Db    682 gcctgccgccggagggacgccctcactggccacctgaggacgcactccgttggtaaacct 741

Qy    601 CACAAATGTGGATATTGTGGCCGAAGCTATAAACAGCGAACGTCTTTAGAGGAACATAAA 660
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    742 cacaaatgtggatattgtggccgaagctataaacagcgaacgtctttagaggaacataaa 801

Qy    661 GAGCGCTGCCACAACTACTTGGAAAGCATGGGCCTTCCGGGCACACTGTACCCAGTCATT 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    802 gagcgctgccacaactacttggaaagcatgggccttccgggcacactgtacccagtcatt 861

Qy    721 AAAGAAGAAACTAAGCACAGTGAAATGGCAGAAGACCTGTGCAAGATAGGATCAGAGAGA 780
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    862 aaagaagaaactaagcacagtgaaatggcagaagacctgtgcaagataggatcagagaga 921

Qy    781 TCTCTCGTGCTGGACAGACTAGCAAGTAATGTCGCCAAACGTAAGAGCTCTATGCCTCAG 840
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    922 tctctcgtgctggacagactagcaagtaatgtcgccaaacgtaagagctctatgcctcag 981

Qy    841 AAATTTCTTGGGGACAAGGGCCTGTCCGACACGCCCTACGACAGTGCCACGTACGAGAAG 900
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    982 aaatttcttggggacaagggcctgtccgacacgccctacgacagtgccacgtacgagaag 1041

Qy    901 GAGAACGAAATGATGAAGTCCCACGTGATGGACCAAGCCATCAACAACGCCATCAACTAC 960
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1042 gagaacgaaatgatgaagtcccacgtgatggaccaagccatcaacaacgccatcaactac 1101

Qy    961 CTGGGGGCCGAGTCCCTGCGCCCGCTGGTGCAGACGCCCCCGGGCGGTTCCGAGGTGGTC 1020
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1102 ctgggggccgagtccctgcgcccgctggtgcagacgccccgggcggttccgaggtggtc 1161

Qy   1021 CCGGTCATCAGCCCGATGTACCAGCTGCACA---GGCGCTCGGAGGGCACCCCGCGCTCC 1077
          |||||||||||||||||||||||||||||||    || | ||||||||||||||||||||
Db   1162 ccggtcatcagcccgatgtaccagctgcacaagccgctcgcggagggcaccccgcgctcc 1221

Qy   1078 AACCACTCGGCCCAGGACAGCGCCGTGGAGTACCTGCTGCTGCTCTCCAAGGCCAAGTTG 1137
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1222 aaccactcggcccaggacagcgccgtggagtacctgctgctgctctccaaggccaagttg 1281

Qy   1138 GTGCCCTCGGAGCGCGAGGCGTCCCCGAGCAACAGCTGCCAAGACTCCACGGACACCGAG 1197
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1282 gtgccctcggagcgcgaggcgtccccgagcaacagctgccaagactccacggacaccgag 1341

Qy   1198 AGCAACAACGAGGAGCAGCGCAGCGGTCTTATCTACCTGACCAACCACATCGCCCGACGC 1257
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||  | |
Db   1342 agcaacaacgaggagcagcgcagcggtcttatctacctgaccaaccacatcgccccgcac 1401

Qy   1258 GCGC---AACGCGTGTCGCTCAAGGAGGAGCACCGCGCCTACGACCTGCTGCGCGCCGCC 1314
          ||||   |  || ||||||||||||||||||||||||||||||||||||||||||||||
Db   1402 gcgcgcaacggcttgtcgctcaaggaggagcaccgcgcctacgacctgctgcgcgccgcc 1461

Qy   1315 TCCGAGAACTCGCAGGACGCGCTCCGCGTGGTCAGCACCAGCGGGGAGCAGATGAAGGTG 1374
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1462 tccgagaactcgcaggacgcgctccgcgtggtcagcaccagcggggagcagatgaaggtg 1521

Qy   1375 TACAAGTGCGAACACTGCCGGGTGCTCTTCCTGGATCACGTCATGTACACCATCCACATG 1434
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1522 tacaagtgcgaacactgccgggtgctcttcctggatcacgtcatgtacaccatccacatg 1581

Qy   1435 GGCTGCCACGGCTTCCGTGATCCTTTTGAGTGCAACATGTGCGGCTACCACAGCCAGGAC 1494
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1582 ggctgccacggcttccgtgatccttttgagtgcaacatgtgcggctaccacagccaggac 1641

Qy   1495 CGGTACGAGTTCTCGTCGCACATAACGCGAGGGGAGCACCGCTTCCACATGAGCTAA 1551
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1642 cggtacgagttctcgtcgcacataacgcgaggggagcaccgcttccacatgagctaa 1698


RESULT   3
US-09-023-655-923
; Sequence 923, Application US/09023655
;  GENERAL INFORMATION:
;    APPLICANT:  Cocks, Benjamin G.
;    APPLICANT:  Susan G. Stuart
```

```
;    APPLICANT:  Jeffrey J. Seilhamer
;    TITLE OF INVENTION:  COMPOSITION FOR THE DETECTION OF BLOOD CELL GENE
;    TITLE OF INVENTION:  EXPRESSION
;    NUMBER OF SEQUENCES:  1508
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  INCYTE PHARMACEUTICALS, INC.
;      STREET:  3174 PORTER DRIVE
;      CITY:  PALO ALTO
;      STATE:  CALIFORNIA
;      COUNTRY:  USA
;      ZIP:  94304
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  Word Perfect 6.1 for Windows/MS-DOS 6.2
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/09/023,655
;      FILING DATE:  HEREWITH
;      CLASSIFICATION:
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:
;      FILING DATE:
;      CLASSIFICATION:
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Zeller, Karen J.
;      REGISTRATION NUMBER:  37,071
;      REFERENCE/DOCKET NUMBER:  PA-0001 US
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (650) 855-0555
;      TELEFAX:  (650) 845-4166
;  INFORMATION FOR SEQ ID NO:  923:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  3629 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  single
;      TOPOLOGY:  linear
;    IMMEDIATE SOURCE:
;      LIBRARY:  GENBANK
;      CLONE:  g1289370
US-09-023-655-923


  Query Match             94.7%;  Score 1468.8;  DB 14;  Length 3629;
  Best Local Similarity   97.7%;  Pred. No. 1.7e-297;
  Matches 1524;  Conservative    0;  Mismatches   27;  Indels    9;  Gaps    3;

Qy       1 ATGGATGCTGACGAGGGTCAAGACATGTCTTTCTCATCAGGGAAGGAAAGCCCCCCTGTA 60
             |||||||||||| ||||||||||||||||||        |||||||||||||||||||||||||||
Db     169 ATGGATGCTGATGAGGGTCAAGACATGTCCCAAGTTTCAGGGAAGGAAAGCCCCCCTGTA 228

Qy      61 AGCGATACTCCAGATGAGGGCGATGAGCCCATGCCGATCCCCGAGGACCTCTCCACCACC 120
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     229 AGCGATACTCCAGATGAGGGCGATGAGCCCATGCCGATCCCCGAGGACCTCTCCACCACC 288

Qy     121 TCGGGAGGACAGCAAAGCTCCAAGAGTGACAGAGTCGTGGCCAGTAATGTTAAAGTAGAG 180
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     289 TCGGGAGGACAGCAAAGCTCCAAGAGTGACAGAGTCGTGGCCAGTAATGTTAAAGTAGAG 348

Qy     181 ACTCAGAGTGATGAAGAGAATGGGCGTGCCTGTGAAATGAATGGGGAAGAATGTGCGGAG 240
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     349 ACTCAGAGTGATGAAGAGAATGGGCGTGCCTGTGAAATGAATGGGGAAGAATGTGCGGAG 408

Qy     241 GATTTACGAATGCTTGATGCCTCGGGAGAGAAAATGAATGGCTCCCACAGGGACCAAGGC 300
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     409 GATTTACGAATGCTTGATGCCTCGGGAGAGAAAATGAATGGCTCCCACAGGGACCAAGGC 468

Qy     301 AGCTCGGCTTTGTCGGGAGTTGGAGGCATTCGACTTCCTAACGGAAAACTAAAGTGTGAT 360
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     469 AGCTCGGCTTTGTCGGGAGTTGGAGGCATTCGACTTCCTAACGGAAAACTAAAGTGTGAT 528

Qy .   361 ATCTGTGGGATCATTTGCATCGGGCCCAATGTGCTCATGGTTCACAAAAGAAGCCACACT 420
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||

Db     529 ATCTGTGGGATCATTTGCATCGGGCCCAATGTGCTCATGGTTCACAAAAGAAGCCACACT 588

Qy     421 GGAGAACGGCCCTTCCAGTGCAATCAGTGCGGGGCCTCATTCACCCAGAAGGGCAACCTG 480
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     589 GGAGAACGGCCCTTCCAGTGCAATCAGTGCGGGGCCTCATTCACCCAGAAGGGCAACCTG 648

Qy     481 CTCCGGCACATCAAGCTGCATTCCGGGGAGAAGCCCTTCAAATGCCACCTCTGCAACTAC 540
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     649 CTCCGGCACATCAAGCTGCATTCCGGGGAGAAGCCCTTCAAATGCCACCTCTGCAACTAC 708

Qy     541 GCCTGCCGCCGGAGGGACGCCCTCACTGGCCACCTGAGGACGCACTCCGTTGGTAAACCT 600
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     709 GCCTGCCGCCGGAGGGACGCCCTCACTGGCCACCTGAGGACGCACTCCGTTGGTAAACCT 768

Qy     601 CACAAATGTGGATATTGTGGCCGAAGCTATAAACAGCGAACGTCTTTAGAGGAACATAAA 660
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     769 CACAAATGTGGATATTGTGGCCGAAGCTATAAACAGCGAAGCTCTTTAGAGGAACATAAA 828

Qy     661 GAGCGCTGCCACAACTACTTGGAAAGCATGGGCCTTCCGGGCACACTGTACCCAGTCATT 720
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     829 GAGCGCTGCCACAACTACTTGGAAAGCATGGGCCTTCCGGGCACACTGTACCCAGTCATT 888

Qy     721 AAAGAAGAAACTAAGCACAGTGAAATGGCAGAAGACCTGTGCAAGATAGGATCAGAGAGA 780
             |||||||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     889 AAAGAAGAAACTAATCACAGTGAAATGGCAGAAGACCTGTGCAAGATAGGATCAGAGAGA 948

Qy     781 TCTCTCGTGCTGGACAGACTAGCAAGTAATGTCGCCAAACGTAAGAGCTCTATGCCTCAG 840
             |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||||||||||||
Db     949 TCTCTCGTGCTGGACAGACTAGCAAGTAACGTCGCCAAACGTAAGAGCTCTATGCCTCAG 1008

Qy     841 AAATTTCTTGGGGACAAGGGCCTGTCCGACACGCCCTACGA---CAGTGCCACGTACGAG 897
             ||||||||||||||||||||||||||||||||||||||||   ||| ||||  ||||||
Db    1009 AAATTTCTTGGGGACAAGGGCCTGTCCGACACGCCCTACGACAGCAGCGCCAGCTACGAG 1068

Qy     898 AAGGAGAACGAAATGATGAAGTCCCACGTGATGGACCAAGCCATCAACAACGCCATCAAC 957
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1069 AAGGAGAACGAAATGATGAAGTCCCACGTGATGGACCAAGCCATCAACAACGCCATCAAC 1128

Qy     958 TACCTGGGGGCCGAGTCCCTGCGCCCGCTGGTGCAGACGCCCCCGGGCGGTTCCGAGGTG 1017
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1129 TACCTGGGGGCCGAGTCCCTGCGCCCGCTGGTGCAGACGCCCCCGGGCGGTTCCGAGGTG 1188

Qy    1018 GTCCCGGTCATCAGCCCGATGTACCAGCTGCACA---GGCGCTCGGAGGGCACCCCGCGC 1074
             ||||||||||||||||||||||||||||||||    || | ||||||||||||||||||
Db    1189 GTCCCGGTCATCAGCCCGATGTACCAGCTGCACAAGCCGCTCGCGGAGGGCACCCCGCGC 1248

Qy    1075 TCCAACCACTCGGCCCAGGACAGCGCCGTGGAGTACCTGCTGCTGCTCTCCAAGGCCAAG 1134
             |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db    1249 TCCAACCACTCGGCCCAGGACAGCGCCGTGGAGAACCTGCTGCTGCTCTCCAAGGCCAAG 1308

Qy    1135 TTGGTGCCCTCGGAGCGCGAGGCGTCCCCGAGCAACAGCTGCCAAGACTCCACGGACACC 1194
             |||||||||||||||||||||||||||||||||||| ||||||||||||||||||||||
Db    1309 TTGGTGCCCTCGGAGCGCGAGGCGTCCCCGAGCAACAGCTGTCAAGACTCCACGGACACC 1368

Qy    1195 GAGAGCAACAACGAGGAGCAGCGCAGCGGTCTTATCTACCTGACCAACCACATCGCCCGA 1254
             |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db.   1369 GAGAGCAACAACGAGGAGCAGCGCAGCGGTCTCATCTACCTGACCAACCACATCGCCCCG 1428

Qy    1255 CGCGCGC---AACGCGTGTCGCTCAAGGAGGAGCACCGCGCCTACGACCTGCTGCGCGCC 1311
             | |||||   |  || ||||||||||||||||||||||||||||||||||||||||||||
Db    1429 CACGCGCGCAACGGCTTGTCGCTCAAGGAGGAGCACCGCGCCTACGACCTGCTGCGCGCC 1488

Qy    1312 GCCTCCGAGAACTCGCAGGACGCGCTCCGCGTGGTCAGCACCAGCGGGGAGCAGATGAAG 1371
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1489 GCCTCCGAGAACTCGCAGGACGCGCTCCGCGTGGTCAGCACCAGCGGGGAGCAGATGAAG 1548

Qy    1372 GTGTACAAGTGCGAACACTGCCGGGTGCTCTTCCTGGATCACGTCATGTACACCATCCAC 1431
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1549 GTGTACAAGTGCGAACACTGCCGGGTGCTCTTCCTGGATCACGTCATGTACACCATCCAC 1608

Qy    1432 ATGGGCTGCCACGGCTTCCGTGATCCTTTTGAGTGCAACATGTGCGGCTACCACAGCCAG 1491
             ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1609 ATGGGCTGCCACGGCTTCCGTGATCCTTTTGAGTGCAACATGTGCGGCTACCACAGCCAG 1668
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Qy   1492 GACCGGTACGAGTTCTCGTCGCACATAACGCGAGGGGAGCACCGCTTCCACATGAGCTAA 1551
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1669 GACCGGTACGAGTTCTCGTCGCACATAACGCGAGGGGAGCACCGCTTCCACATGAGCTAA 1728
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RESULT 4
US-09-435-327A-18
; Sequence 18, Application US/09435327A
; GENERAL INFORMATION:
; APPLICANT: Uckun, Fatih M.
; APPLICANT: Crotty, Mya L.
; TITLE OF INVENTION: IKAROS ISOFORMS AND MUTANTS
; FILE REFERENCE: 12152.35USU1
; CURRENT APPLICATION NUMBER: US/09/435,327A
; CURRENT FILING DATE: 1999-11-05
; PRIOR APPLICATION NUMBER: 60/107,229
; PRIOR FILING DATE: 1998-11-05
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 18
; LENGTH: 3629
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (3078)
; OTHER INFORMATION: Genbank HSU40462
US-09-435-327A-18

Query Match 94.7%; Score 1468.8; DB 18; Length 3629;
Best Local Similarity 97.7%; Pred. No. 1.7e-297;
Matches 1524; Conservative 0; Mismatches 27; Indels 9; Gaps 3;

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Qy      1 ATGGATGCTGACGAGGGTCAAGACATGTCTTTCTCATCAGGGAAGGAAAGCCCCCCTGTA 60
          |||||||||| ||||||||||||||||       |||||||||||||||||||||||||
Db    169 atggatgctgatgagggtcaagacatgtcccaagtttcagggaaggaaagccccccctgta 228

Qy     61 AGCGATACTCCAGATGAGGGCGATGAGCCCATGCCGATCCCCGAGGACCTCTCCACCACC 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    229 agcgatactccagatgagggcgatgagcccatgccgatccccgaggacctctccaccacc 288

Qy    121 TCGGGAGGACAGCAAAGCTCCAAGAGTGACAGAGTCGTGGCCAGTAATGTTAAAGTAGAG 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    289 tcgggaggacagcaaagctccaagagtgacagagtcgtggccagtaatgttaaagtagag 348

Qy    181 ACTCAGAGTGATGAAGAGAATGGGCGTGCCTGTGAAATGAATGGGGAAGAATGTGCGGAG 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    349 actcagagtgatgaagagaatgggcgtgcctgtgaaatgaatggggaagaatgtgcggag 408

Qy    241 GATTTACGAATGCTTGATGCCTCGGGAGAGAAAATGAATGGCTCCCACAGGGACCAAGGC 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    409 gatttacgaatgcttgatgcctcgggagagaaaatgaatggctcccacagggaccaaggc 468

Qy    301 AGCTCGGCTTTGTCGGGAGTTGGAGGCATTCGACTTCCTAACGGAAAACTAAAGTGTGAT 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    469 agctcggctttgtcgggagttggaggcattcgacttcctaacggaaaactaaagtgtgat 528

Qy    361 ATCTGTGGGATCATTTGCATCGGGCCCAATGTGCTCATGGTTCACAAAAGAAGCCACACT 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    529 atctgtgggatcatttgcatcgggcccaatgtgctcatggttcacaaaagaagccacact 588

Qy    421 GGAGAACGGCCCTTCCAGTGCAATCAGTGCGGGGCCTCATTCACCCAGAAGGGCAACCTG 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    589 ggagaacggcccttccagtgcaatcagtgcggggcctcattcacccagaagggcaacctg 648

Qy    481 CTCCGGCACATCAAGCTGCATTCCGGGGAGAAGCCCTTCAAATGCCACCTCTGCAACTAC 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    649 ctccggcacatcaagctgcattccggggagaagcccttcaaatgccacctctgcaactac 708

Qy    541 GCCTGCCGCCGGAGGGACGCCCTCACTGGCCACCTGAGGACGCACTCCGTTGGTAAACCT 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||

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Db   709 gcctgccgccggagggacgccctcactggccacctgaggacgcactccgttggtaaacct 768

Qy   601 CACAAATGTGGATATTGTGGCCGAAGCTATAAACAGCGAACGTCTTTAGAGGAACATAAA 660
          ||||||||||||||||||||||||||||||||||||||||||| ||||||||||||||||||
Db   769 cacaaatgtggatattgtggccgaagctataaacagcgaagctctttagaggaacataaa 828

Qy   661 GAGCGCTGCCACAACTACTTGGAAAGCATGGGCCTTCCGGGCACACTGTACCCAGTCATT 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   829 gagcgctgccacaactacttggaaagcatgggccttccgggcacactgtacccagtcatt 888

Qy   721 AAAGAAGAAACTAAGCACAGTGAAATGGCAGAAGACCTGTGCAAGATAGGATCAGAGAGA 780
          ||||||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||
Db   889 aaagaagaaactaatcacagtgaaatggcagaagacctgtgcaagataggatcagagaga 948

Qy   781 TCTCTCGTGCTGGACAGACTAGCAAGTAATGTCGCCAAACGTAAGAGCTCTATGCCTCAG 840
          |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||||
Db   949 tctctcgtgctggacagactagcaagtaacgtcgccaaacgtaagagctctatgcctcag 1008

Qy   841 AAATTTCTTGGGGACAAGGGCCTGTCCGACACGCCCTACGA---CAGTGCCACGTACGAG 897
          ||||||||||||||||||||||||||||||||||||||||| ||| |||| ||||||
Db   1009 aaatttcttggggacaagggcctgtccgacacgccctacgacagcagcgccagctacgag 1068

Qy   898 AAGGAGAACGAAATGATGAAGTCCCACGTGATGGACCAAGCCATCAACAACGCCATCAAC 957
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1069 aaggagaacgaaatgatgaagtcccacgtgatggaccaagccatcaacaacgccatcaac 1128

Qy   958 TACCTGGGGGCCGAGTCCCTGCGCCCGCTGGTGCAGACGCCCCCGGGCGGTTCCGAGGTG 1017
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1129 tacctgggggccgagtccctgcgcccgctggtgcagacgcccccgggcggttccgaggtg 1188

Qy   1018 GTCCCGGTCATCAGCCCGATGTACCAGCTGCACA---GGCGCTCGGAGGGCACCCCGCGC 1074
          ||||||||||||||||||||||||||||||||| || | ||||||||||||||||
Db   1189 gtcccggtcatcagcccgatgtaccagctgcacaagccgctcgcggagggcaccccgcgc 1248

Qy   1075 TCCAACCACTCGGCCCAGGACAGCGCCGTGGAGTACCTGCTGCTGCTCTCCAAGGCCAAG 1134
          |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db   1249 tccaaccactcggcccaggacagcgccgtggagaacctgctgctgctctccaaggccaag 1308

Qy   1135 TTGGTGCCCTCGGAGCGCGAGGCGTCCCCGAGCAACAGCTGCCAAGACTCCACGGACACC 1194
          |||||||||||||||||||||||||||||||||||||||||||| ||||||||||||||||||
Db   1309 ttggtgccctcggagcgcgaggcgtccccgagcaacagctgtcaagactccacggacacc 1368

Qy   1195 GAGAGCAACAACGAGGAGCAGCGCAGCGGTCTTATCTACCTGACCAACCACATCGCCCGA 1254
          |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db   1369 gagagcaacaacgaggagcagcgcagcggtctcatctacctgaccaaccacatcgccccg 1428

Qy   1255 CGCGCGC---AACGCGTGTCGCTCAAGGAGGAGCACCGCGCCTACGACCTGCTGCGCGCC 1311
          | ||||| | || ||||||||||||||||||||||||||||||||||||||||||||
Db   1429 cacgcgcgcaacggcttgtcgctcaaggaggagcaccgcgcctacgacctgctgcgcgcc 1488

Qy   1312 GCCTCCGAGAACTCGCAGGACGCGCTCCGCGTGGTCAGCACCAGCGGGGAGCAGATGAAG 1371
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1489 gcctccgagaactcgcaggacgcgctccgcgtggtcagcaccagcggggagcagatgaag 1548

Qy   1372 GTGTACAAGTGCGAACACTGCCGGGTGCTCTTCCTGGATCACGTCATGTACACCATCCAC 1431
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1549 gtgtacaagtgcgaacactgccgggtgctcttcctggatcacgtcatgtacaccatccac 1608

Qy   1432 ATGGGCTGCCACGGCTTCCGTGATCCTTTTGAGTGCAACATGTGCGGCTACCACAGCCAG 1491
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1609 atgggctgccacggcttccgtgatccttttgagtgcaacatgtgcggctaccacagccag 1668

Qy   1492 GACCGGTACGAGTTCTCGTCGCACATAACGCGAGGGGAGCACCGCTTCCACATGAGCTAA 1551
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1669 gaccggtacgagttctcgtcgcacataacgcgaggggagcaccgcttccacatgagctaa 1728

RESULT 5
US-09-442-384A-721
; Sequence 721, Application US/09442384A
; GENERAL INFORMATION:
; APPLICANT: Chenchik, Alex
; APPLICANT: Lukashev, Matvey

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;  TITLE OF INVENTION: Hematology/Immunology Array
;  FILE REFERENCE: CLON-006CIP15
;  CURRENT APPLICATION NUMBER: US/09/442,384A
;  CURRENT FILING DATE:  1999-11-17
;  PRIOR APPLICATION NUMBER: 09/053,375
;  PRIOR FILING DATE: 1998-03-31
;  NUMBER OF SEQ ID NOS: 830
;  SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 721
;   LENGTH: 3629
;   TYPE: DNA
;   ORGANISM: homo sapiens
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: 3078
;   OTHER INFORMATION: n = A,T,C or G
;   NAME/KEY: misc_feature
;   LOCATION: 3078
;   OTHER INFORMATION: n = A,T,C or G
US-09-442-384A-721

  Query Match             94.7%;  Score 1468.8;  DB 18;  Length 3629;
  Best Local Similarity   97.7%;  Pred. No. 1.7e-297;
  Matches 1524;  Conservative    0;  Mismatches   27;  Indels    9;  Gaps    3;

Qy       1 ATGGATGCTGACGAGGGTCAAGACATGTCTTTCTCATCAGGGAAGGAAAGCCCCCCTGTA 60
           |||||||||||| ||||||||||||||||       |||||||||||||||||||||||||
Db     169 atggatgctgatgagggtcaagacatgtcccaagtttcagggaaggaaagccccctgta 228

Qy      61 AGCGATACTCCAGATGAGGGCGATGAGCCCATGCCGATCCCCGAGGACCTCTCCACCACC 120
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     229 agcgatactccagatgagggcgatgagcccatgccgatccccgaggacctctccaccacc 288

Qy     121 TCGGGAGGACAGCAAAGCTCCAAGAGTGACAGAGTCGTGGCCAGTAATGTTAAAGTAGAG 180
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     289 tcgggaggacagcaaagctccaagagtgacagagtcgtggccagtaatgttaaagtagag 348

Qy     181 ACTCAGAGTGATGAAGAGAATGGGCGTGCCTGTGAAATGAATGGGGAAGAATGTGCGGAG 240
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     349 actcagagtgatgaagagaatgggcgtgcctgtgaaatgaatggggaagaatgtgcggag 408

Qy     241 GATTTACGAATGCTTGATGCCTCGGGAGAGAAAATGAATGGCTCCCACAGGGACCAAGGC 300
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     409 gatttacgaatgcttgatgcctcgggagagaaaatgaatggctcccacagggaccaaggc 468

Qy     301 AGCTCGGCTTTGTCGGGAGTTGGAGGCATTCGACTTCCTAACGGAAAACTAAAGTGTGAT 360
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     469 agctcggctttgtcgggagttggaggcattcgacttcctaacggaaaactaaagtgtgat 528

Qy     361 ATCTGTGGGATCATTTGCATCGGGCCCAATGTGCTCATGGTTCACAAAAGAAGCCACACT 420
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     529 atctgtgggatcatttgcatcgggcccaatgtgctcatggttcacaaaagaagccacact 588

Qy     421 GGAGAACGGCCCTTCCAGTGCAATCAGTGCGGGGCCTCATTCACCCAGAAGGGCAACCTG 480
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     589 ggagaacggcccttccagtgcaatcagtgcggggcctcattcacccagaagggcaacctg 648

Qy     481 CTCCGGCACATCAAGCTGCATTCCGGGGAGAAGCCCTTCAAATGCCACCTCTGCAACTAC 540
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     649 ctccggcacatcaagctgcattccggggagaagcccttcaaatgccacctctgcaactac 708

Qy     541 GCCTGCCGCCGGAGGGACGCCCTCACTGGCCACCTGAGGACGCACTCCGTTGGTAAACCT 600
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     709 gcctgccgccggagggacgccctcactggccacctgaggacgcactccgttggtaaacct 768

Qy     601 CACAAATGTGGATATTGTGGCCGAAGCTATAAACAGCGAACGTCTTTAGAGGAACATAAA 660
           |||||||||||||||||||||||||||||||||||  ||||||||||||||||||||||
Db     769 cacaaatgtggatattgtggccgaagctataaacagcgaagctctttagaggaacataaa 828

Qy     661 GAGCGCTGCCACAACTACTTGGAAAGCATGGGCCTTCCGGGCACACTGTACCCAGTCATT 720
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     829 gagcgctgccacaactacttggaaagcatgggccttccgggcacactgtacccagtcatt 888

Qy     721 AAAGAAGAAACTAAGCACAGTGAAATGGCAGAAGACCTGTGCAAGATAGGATCAGAGAGA 780
           |||||||||||||| ||||||||||||||||||||||||||||||||||||||||||||||
Db     889 aaagaagaaactaatcacagtgaaatggcagaagacctgtgcaagataggatcagagaga 948

Qy     781 TCTCTCGTGCTGGACAGACTAGCAAGTAATGTCGCCAAACGTAAGAGCTCTATGCCTCAG 840
           |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||||
Db     949 tctctcgtgctggacagactagcaagtaacgtcgccaaacgtaagagctctatgcctcag 1008

Qy     841 AAATTTCTTGGGGACAAGGGCCTGTCCGACACGCCCTACGA---CAGTGCCACGTACGAG 897
           |||||||||||||||||||||||||||||||||||||||||   ||| ||||  ||||||
Db    1009 aaatttcttggggacaagggcctgtccgacacgccctacgacagcagcgccagctacgag 1068

Qy     898 AAGGAGAACGAAATGATGAAGTCCCACGTGATGGACCAAGCCATCAACAACGCCATCAAC 957
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1069 aaggagaacgaaatgatgaagtcccacgtgatggaccaagccatcaacaacgccatcaac 1128

Qy     958 TACCTGGGGGCCGAGTCCCTGCGCCCGCTGGTGCAGACGCCCCCGGGCGGTTCCGAGGTG 1017
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1129 tacctgggggccgagtccctgcgcccgctggtgcagacgccccgggcggttccgaggtg 1188

Qy    1018 GTCCCGGTCATCAGCCCGATGTACCAGCTGCACA---GGCGCTCGGAGGGCACCCCGCGC 1074
           |||||||||||||||||||||||||||||||||     || | ||||||||||||||||
Db    1189 gtcccggtcatcagcccgatgtaccagctgcacaagccgctcgcggagggcaccccgcgc 1248

Qy    1075 TCCAACCACTCGGCCCAGGACAGCGCCGTGGAGTACCTGCTGCTGCTCTCCAAGGCCAAG 1134
           |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db    1249 tccaaccactcggcccaggacagcgccgtggagaacctgctgctgctctccaaggccaag 1308

Qy    1135 TTGGTGCCCTCGGAGCGCGAGGCGTCCCCGAGCAACAGCTGCCAAGACTCCACGGACACC 1194
           |||||||||||||||||||||||||||||||||||||||||||||| ||||||||||||||
Db    1309 ttggtgccctcggagcgcgaggcgtccccgagcaacagctgtcaagactccacggacacc 1368

Qy    1195 GAGAGCAACAACGAGGAGCAGCGCAGCGGTCTTATCTACCTGACCAACCACATCGCCCGA 1254
           |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db    1369 gagagcaacaacgaggagcagcgcagcggtctcatctacctgaccaaccacatcgccccg 1428

Qy    1255 CGCGCGC---AACGCGTGTCGCTCAAGGAGGAGCACCGCGCCTACGACCTGCTGCGCGCC 1311
           | |||||   |  || ||||||||||||||||||||||||||||||||||||||||||||
Db    1429 cacgcgcgcaacggcttgtcgctcaaggaggagcaccgcgcctacgacctgctgcgcgcc 1488

Qy    1312 GCCTCCGAGAACTCGCAGGACGCGCTCCGCGTGGTCAGCACCAGCGGGGAGCAGATGAAG 1371
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1489 gcctccgagaactcgcaggacgcgctccgcgtggtcagcaccagcggggagcagatgaag 1548

Qy    1372 GTGTACAAGTGCGAACACTGCCGGGTGCTCTTCCTGGATCACGTCATGTACACCATCCAC 1431
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1549 gtgtacaagtgcgaacactgccgggtgctcttcctggatcacgtcatgtacaccatccac 1608

Qy    1432 ATGGGCTGCCACGGCTTCCGTGATCCTTTTGAGTGCAACATGTGCGGCTACCACAGCCAG 1491
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1609 atgggctgccacggcttccgtgatccttttgagtgcaacatgtgcggctaccacagccag 1668

Qy    1492 GACCGGTACGAGTTCTCGTCGCACATAACGCGAGGGGAGCACCGCTTCCACATGAGCTAA 1551
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1669 gaccggtacgagttctcgtcgcacataacgcgaggggagcaccgcttccacatgagctaa 1728

RESULT   6
US-09-649-161-10565
; Sequence 10565, Application US/09649161
; GENERAL INFORMATION:
;  APPLICANT:  Gutierrez-Ramos, Jose-Carlos
;  APPLICANT:  Villeval, Jean-Luc M.G.
;  APPLICANT:  Fraser, Christopher C.
;  APPLICANT:  Holtzman, Douglas A.
;  APPLICANT:  Hodge, Martin R.
;  TITLE OF INVENTION:  NOVEL NUCLEIC ACID MOLECULES AND USES
;  TITLE OF INVENTION:  THEREFOR
;  FILE REFERENCE:  1600.1178-001
;  CURRENT APPLICATION NUMBER: US/09/649,161
;  CURRENT FILING DATE:  2000-08-28
;  PRIOR APPLICATION NUMBER:  60/151,060
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;  PRIOR FILING DATE:  1999-08-27
;  NUMBER OF SEQ ID NOS:  10788
;  SOFTWARE:  FastSEQ for Windows Version 4.0
; SEQ ID NO 10565
;   LENGTH: 4786
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(4786)
;   OTHER INFORMATION: n = A,T,C or G
US-09-649-161-10565
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Query Match 94.7%; Score 1468.8; DB 25; Length 4786;
Best Local Similarity 97.7%; Pred. No. 1.8e-297;
Matches 1524; Conservative 0; Mismatches 27; Indels 9; Gaps 3;

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Qy       1 ATGGATGCTGACGAGGGTCAAGACATGTCTTTCTCATCAGGGAAGGAAAGCCCCCCTGTA 60
          ||||||||||| |||||||||||||||||       |||||||||||||||||||||||||
Db     434 atggatgctgatgagggtcaagacatgtcccaagtttcagggaaggaaagccccctgta 493

Qy      61 AGCGATACTCCAGATGAGGGCGATGAGCCCATGCCGATCCCCGAGGACCTCTCCACCACC 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     494 agcgatactccagatgagggcgatgagcccatgccgatccccgaggacctctccaccacc 553

Qy     121 TCGGGAGGACAGCAAAGCTCCAAGAGTGACAGAGTCGTGGCCAGTAATGTTAAAGTAGAG 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     554 tcgggaggacagcaaagctccaagagtgacagagtcgtggccagtaatgttaaagtagag 613

Qy     181 ACTCAGAGTGATGAAGAGAATGGGCGTGCCTGTGAAATGAATGGGGAAGAATGTGCGGAG 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     614 actcagagtgatgaagagaatgggcgtgcctgtgaaatgaatggggaagaatgtgcggag 673

Qy     241 GATTTACGAATGCTTGATGCCTCGGGAGAGAAAATGAATGGCTCCCACAGGGACCAAGGC 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     674 gatttacgaatgcttgatgcctcgggagagaaaatgaatggctcccacagggaccaaggc 733

Qy     301 AGCTCGGCTTTGTCGGGAGTTGGAGGCATTCGACTTCCTAACGGAAAACTAAAGTGTGAT 360
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     734 agctcggctttgtcgggagttggaggcattcgacttcctaacggaaaactaaagtgtgat 793

Qy     361 ATCTGTGGGATCATTTGCATCGGGCCCAATGTGCTCATGGTTCACAAAAGAAGCCACACT 420
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     794 atctgtgggatcatttgcatcgggcccaatgtgctcatggttcacaaaagaagccacact 853

Qy     421 GGAGAACGGCCCTTCCAGTGCAATCAGTGCGGGGCCTCATTCACCCAGAAGGGCAACCTG 480
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     854 ggagaacggcccttccagtgcaatcagtgcggggcctcattcacccagaagggcaacctg 913

Qy     481 CTCCGGCACATCAAGCTGCATTCCGGGGAGAAGCCCTTCAAATGCCACCTCTGCAACTAC 540
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     914 ctccggcacatcaagctgcattccggggagaagcccttcaaatgccacctctgcaactac 973

Qy     541 GCCTGCCGCCGGAGGGACGCCCTCACTGGCCACCTGAGGACGCACTCCGTTGGTAAACCT 600
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     974 gcctgccgccggagggacgccctcactggccacctgaggacgcactccgttggtaaacct 1033

Qy     601 CACAAATGTGGATATTGTGGCCGAAGCTATAAACAGCGAACGTCTTTAGAGGAACATAAA 660
          ||||||||||||||||||||||||||||||||||||  ||||||||||||||||||||
Db    1034 cacaaatgtggatattgtggccgaagctataaacagcgaagctctttagaggaacataaa 1093

Qy     661 GAGCGCTGCCACAACTACTTGGAAAGCATGGGCCTTCCGGGCACACTGTACCCAGTCATT 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1094 gagcgctgccacaactacttggaaagcatgggccttccgggcacactgtacccagtcatt 1153

Qy     721 AAAGAAGAAACTAAGCACAGTGAAATGGCAGAAGACCTGTGCAAGATAGGATCAGAGAGA 780
          |||||||||||||| ||||||||||||||||||||||||||||||||||||||||||||
Db    1154 aaagaagaaactaatcacagtgaaatggcagaagacctgtgcaagataggatcagagaga 1213

Qy     781 TCTCTCGTGCTGGACAGACTAGCAAGTAATGTCGCCAAACGTAAGAGCTCTATGCCTCAG 840
          |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db    1214 tctctcgtgctggacagactagcaagtaacgtcgccaaacgtaagagctctatgcctcag 1273

Qy     841 AAATTTCTTGGGGACAAGGGCCTGTCCGACACGCCCTACGA---CAGTGCCACGTACGAG 897
          |||||||||||||||||||||||||||||||||||||||||   ||| |||| ||||||
Db    1274 aaatttcttggggacaagggcctgtccgacacgccctacgacagcagcgccagctacgag 1333

Qy     898 AAGGAGAACGAAATGATGAAGTCCCACGTGATGGACCAAGCCATCAACAACGCCATCAAC 957
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1334 aaggagaacgaaatgatgaagtcccacgtgatggaccaagccatcaacaacgccatcaac 1393

Qy     958 TACCTGGGGGCCGAGTCCCTGCGCCCGCTGGTGCAGACGCCCCCGGGCGGTTCCGAGGTG 1017
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1394 tacctgggggccgagtccctgcgcccgctggtgcagacgccccgggcggttccgaggtg 1453

Qy    1018 GTCCCGGTCATCAGCCCGATGTACCAGCTGCACA---GGCGCTCGGAGGGCACCCCGCGC 1074
          ||||||||||||||||||||||||||||||||    || | ||||||||||||||||||
Db    1454 gtcccggtcatcagcccgatgtaccagctgcacaagccgctcgcggagggcaccccgcgc 1513

Qy    1075 TCCAACCACTCGGCCCAGGACAGCGCCGTGGAGTACCTGCTGCTGCTCTCCAAGGCCAAG 1134
          |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db    1514 tccaaccactcggcccaggacagcgccgtggagaacctgctgctgctctccaaggccaag 1573

Qy    1135 TTGGTGCCCTCGGAGCGCGAGGCGTCCCCGAGCAACAGCTGCCAAGACTCCACGGACACC 1194
          |||||||||||||||||||||||||||||||||||||||||| ||||||||||||||||||
Db    1574 ttggtgccctcggagcgcgaggcgtccccgagcaacagctgtcaagactccacggacacc 1633

Qy    1195 GAGAGCAACAACGAGGAGCAGCGCAGCGGTCTTATCTACCTGACCAACCACATCGCCCGA 1254
          |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db    1634 gagagcaacaacgaggagcagcgcagcggtctcatctacctgaccaaccacatcgccccg 1693

Qy    1255 CGCGCGC---AACGCGTGTCGCTCAAGGAGGAGCACCGCGCCTACGACCTGCTGCGCGCC 1311
          | |||||   |  || ||||||||||||||||||||||||||||||||||||||||||||
Db    1694 cacgcgcgcaacggcttgtcgctcaaggaggagcaccgcgcctacgacctgctgcgcgcc 1753

Qy    1312 GCCTCCGAGAACTCGCAGGACGCGCTCCGCGTGGTCAGCACCAGCGGGGAGCAGATGAAG 1371
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1754 gcctccgagaactcgcaggacgcgctccgcgtggtcagcaccagcggggagcagatgaag 1813

Qy    1372 GTGTACAAGTGCGAACACTGCCGGGTGCTCTTCCTGGATCACGTCATGTACACCATCCAC 1431
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1814 gtgtacaagtgcgaacactgccgggtgctcttcctggatcacgtcatgtacaccatccac 1873

Qy    1432 ATGGGCTGCCACGGCTTCCGTGATCCTTTTGAGTGCAACATGTGCGGCTACCACAGCCAG 1491
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1874 atgggctgccacggcttccgtgatccttttgagtgcaacatgtgcggctaccacagccag 1933

Qy    1492 GACCGGTACGAGTTCTCGTCGCACATAACGCGAGGGGAGCACCGCTTCCACATGAGCTAA 1551
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1934 gaccggtacgagttctcgtcgcacataacgcgaggggagcaccgcttccacatgagctaa 1993
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RESULT   7
US-09-652-816-9158
; Sequence 9158, Application US/09652816
; GENERAL INFORMATION:
;  APPLICANT:  Gutierrez-Ramos, Jose-Carlos
;  TITLE OF INVENTION:  NOVEL NUCLEIC ACID MOLECULES AND USES
;  TITLE OF INVENTION:  THEREFOR
;  FILE REFERENCE:  1600.1177-001
;  CURRENT APPLICATION NUMBER: US/09/652,816
;  CURRENT FILING DATE:  2000-08-31
;  PRIOR APPLICATION NUMBER:  60/152,111
;  PRIOR FILING DATE:  1999-08-31
;  NUMBER OF SEQ ID NOS:  9647
;  SOFTWARE:  FastSEQ for Windows Version 4.0
; SEQ ID NO 9158
;   LENGTH: 4786
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(4786)
;   OTHER INFORMATION: n = A,T,C or G
US-09-652-816-9158
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  Query Match             94.7%;  Score 1468.8;  DB 25;  Length 4786;
  Best Local Similarity   97.7%;  Pred. No. 1.8e-297;
  Matches 1524;  Conservative    0;  Mismatches   27;  Indels    9;  Gaps    3;

Qy       1 ATGGATGCTGACGAGGGTCAAGACATGTCTTTCTCATCAGGGAAGGAAAGCCCCCCTGTA 60
           |||||||||| |||||||||||||||       |||||||||||||||||||||||||
Db     434 atggatgctgatgagggtcaagacatgtcccaagtttcagggaaggaaagccccccctgta 493

Qy      61 AGCGATACTCCAGATGAGGGCGATGAGCCCATGCCGATCCCCGAGGACCTCTCCACCACC 120
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     494 agcgatactccagatgagggcgatgagcccatgccgatccccgaggacctctccaccacc 553

Qy     121 TCGGGAGGACAGCAAAGCTCCAAGAGTGACAGAGTCGTGGCCAGTAATGTTAAAGTAGAG 180
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     554 tcgggaggacagcaaagctccaagagtgacagagtcgtggccagtaatgttaaagtagag 613

Qy     181 ACTCAGAGTGATGAAGAGAATGGGCGTGCCTGTGAAATGAATGGGGAAGAATGTGCGGAG 240
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     614 actcagagtgatgaagagaatgggcgtgcctgtgaaatgaatggggaagaatgtgcggag 673

Qy     241 GATTTACGAATGCTTGATGCCTCGGGAGAGAAAATGAATGGCTCCCACAGGGACCAAGGC 300
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     674 gatttacgaatgcttgatgcctcgggagagaaaatgaatggctcccacagggaccaaggc 733

Qy     301 AGCTCGGCTTTGTCGGGAGTTGGAGGCATTCGACTTCCTAACGGAAAACTAAAGTGTGAT 360
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     734 agctcggctttgtcgggagttggaggcattcgacttcctaacggaaaactaaagtgtgat 793

Qy     361 ATCTGTGGGATCATTTGCATCGGGCCCAATGTGCTCATGGTTCACAAAAGAAGCCACACT 420
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     794 atctgtgggatcatttgcatcgggcccaatgtgctcatggttcacaaaagaagccacact 853

Qy     421 GGAGAACGGCCCTTCCAGTGCAATCAGTGCGGGGCCTCATTCACCCAGAAGGGCAACCTG 480
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     854 ggagaacggcccttccagtgcaatcagtgcggggcctcattcacccagaagggcaacctg 913

Qy     481 CTCCGGCACATCAAGCTGCATTCCGGGGAGAAGCCCTTCAAATGCCACCTCTGCAACTAC 540
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     914 ctccggcacatcaagctgcattccggggagaagcccttcaaatgccacctctgcaactac 973

Qy     541 GCCTGCCGCCGGAGGGACGCCCTCACTGGCCACCTGAGGACGCACTCCGTTGGTAAACCT 600
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     974 gcctgccgccggagggacgccctcactggccacctgaggacgcactccgttggtaaacct 1033

Qy     601 CACAAATGTGGATATTGTGGCCGAAGCTATAAACAGCGAACGTCTTTAGAGGAACATAAA 660
           ||||||||||||||||||||||||||||||||||||  ||||||||||||||||||||
Db    1034 cacaaatgtggatattgtggccgaagctataaacagcgaagctctttagaggaacataaa 1093

Qy     661 GAGCGCTGCCACAACTACTTGGAAAGCATGGGCCTTCCGGGCACACTGTACCCAGTCATT 720
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1094 gagcgctgccacaactacttggaaagcatgggccttccgggcacactgtacccagtcatt 1153

Qy     721 AAAGAAGAAACTAAGCACAGTGAAATGGCAGAAGACCTGTGCAAGATAGGATCAGAGAGA 780
           |||||||||||||| ||||||||||||||||||||||||||||||||||||||||||||
Db    1154 aaagaagaaactaatcacagtgaaatggcagaagacctgtgcaagataggatcagagaga 1213

Qy     781 TCTCTCGTGCTGGACAGACTAGCAAGTAATGTCGCCAAACGTAAGAGCTCTATGCCTCAG 840
           |||||||||||||||||||||||||||| ||||||||||||||||||||||||||||||
Db    1214 tctctcgtgctggacagactagcaagtaacgtcgccaaacgtaagagctctatgcctcag 1273

Qy     841 AAATTTCTTGGGGACAAGGGCCTGTCCGACACGCCCTACGA---CAGTGCCACGTACGAG 897
           |||||||||||||||||||||||||||||||||||||||||   ||| ||||  ||||||
Db    1274 aaatttcttggggacaagggcctgtccgacacgccctacgacagcagcgccagctacgag 1333

Qy     898 AAGGAGAACGAAATGATGAAGTCCCACGTGATGGACCAAGCCATCAACAACGCCATCAAC 957
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1334 aaggagaacgaaatgatgaagtcccacgtgatggaccaagccatcaacaacgccatcaac 1393

Qy     958 TACCTGGGGGCCGAGTCCCTGCGCCCGCTGGTGCAGACGCCCCCGGGCGGTTCCGAGGTG 1017
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1394 tacctgggggccgagtccctgcgcccgctggtgcagacgccccgggcggttccgaggtg 1453

Qy    1018 GTCCCGGTCATCAGCCCGATGTACCAGCTGCACA---GGCGCTCGGAGGGCACCCCGCGC 1074
           ||||||||||||||||||||||||||||||||    || | ||||||||||||||||
Db    1454 gtcccggtcatcagcccgatgtaccagctgcacaagccgctcgcggagggcaccccgcgc 1513

Qy    1075 TCCAACCACTCGGCCCAGGACAGCGCCGTGGAGTACCTGCTGCTGCTCTCCAAGGCCAAG 1134
           |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db    1514 tccaaccactcggcccaggacagcgccgtggagaacctgctgctgctctccaaggccaag 1573

Qy    1135 TTGGTGCCCTCGGAGCGCGAGGCGTCCCCGAGCAACAGCTGCCAAGACTCCACGGACACC 1194
           |||||||||||||||||||||||||||||||||||||||||| ||||||||||||||||
Db    1574 ttggtgccctcggagcgcgaggcgtccccgagcaacagctgtcaagactccacggacacc 1633

Qy    1195 GAGAGCAACAACGAGGAGCAGCGCAGCGGTCTTATCTACCTGACCAACCACATCGCCCGA 1254
           |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db    1634 gagagcaacaacgaggagcagcgcagcggtctcatctacctgaccaaccacatcgccccg 1693

Qy    1255 CGCGCGC---AACGCGTGTCGCTCAAGGAGGAGCACCGCGCCTACGACCTGCTGCGCGCC 1311
           | |||||   | || ||||||||||||||||||||||||||||||||||||||||||
Db    1694 cacgcgcgcaacggcttgtcgctcaaggaggagcaccgcgcctacgacctgctgcgcgcc 1753

Qy    1312 GCCTCCGAGAACTCGCAGGACGCGCTCCGCGTGGTCAGCACCAGCGGGGAGCAGATGAAG 1371
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1754 gcctccgagaactcgcaggacgcgctccgcgtggtcagcaccagcggggagcagatgaag 1813

Qy    1372 GTGTACAAGTGCGAACACTGCCGGGTGCTCTTCCTGGATCACGTCATGTACACCATCCAC 1431
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1814 gtgtacaagtgcgaacactgccgggtgctcttcctggatcacgtcatgtacaccatccac 1873

Qy    1432 ATGGGCTGCCACGGCTTCCGTGATCCTTTTGAGTGCAACATGTGCGGCTACCACAGCCAG 1491
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1874 atgggctgccacggcttccgtgatcctttgagtgcaacatgtgcggctaccacagccag 1933

Qy    1492 GACCGGTACGAGTTCTCGTCGCACATAACGCGAGGGGAGCACCGCTTCCACATGAGCTAA 1551
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1934 gaccggtacgagttctcgtcgcacataacgcgaggggagcaccgcttccacatgagctaa 1993


RESULT   8
US-09-698-013-6274
; Sequence 6274, Application US/09698013
; GENERAL INFORMATION:
;  APPLICANT:  Gearing, David P.
;  APPLICANT:  Comrack, Christopher
;  APPLICANT:  Kingsbury, Gillian A.
;  APPLICANT:  Holtzman, Douglas A.
;  TITLE OF INVENTION:  NOVEL NUCLEIC ACID MOLECULES AND USES
;  TITLE OF INVENTION:  THEREFOR
;  FILE REFERENCE:  1600.2013-001
;  CURRENT APPLICATION NUMBER: US/09/698,013
;  CURRENT FILING DATE:  2000-10-27
;  PRIOR APPLICATION NUMBER:  60/162,360
;  PRIOR FILING DATE:  1999-10-29
;  NUMBER OF SEQ ID NOS:  7935
;  SOFTWARE:  FastSEQ for Windows Version 4.0
; SEQ ID NO 6274
;   LENGTH: 4786
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(4786)
;   OTHER INFORMATION: n = A,T,C or G
US-09-698-013-6274


  Query Match             94.7%;  Score 1468.8;  DB 27;  Length 4786;
  Best Local Similarity   97.7%;  Pred. No. 1.8e-297;
  Matches 1524;  Conservative    0;  Mismatches   27;  Indels    9;  Gaps    3;

Qy       1 ATGGATGCTGACGAGGGTCAAGACATGTCTTTCTCATCAGGGAAGGAAAGCCCCCCTGTA 60
           |||||||||| |||||||||||||||       |||||||||||||||||||||||||
Db     434 atggatgctgatgagggtcaagacatgtcccaagtttcagggaaggaaagccccccctgta 493
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Qy      61 AGCGATACTCCAGATGAGGGCGATGAGCCCATGCCGATCCCCGAGGACCTCTCCACCACC 120
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     494 agcgatactccagatgagggcgatgagcccatgccgatccccgaggacctctccaccacc 553

Qy     121 TCGGGAGGACAGCAAAGCTCCAAGAGTGACAGAGTCGTGGCCAGTAATGTTAAAGTAGAG 180
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     554 tcgggaggacagcaaagctccaagagtgacagagtcgtggccagtaatgttaaagtagag 613

Qy     181 ACTCAGAGTGATGAAGAGAATGGGCGTGCCTGTGAAATGAATGGGGAAGAATGTGCGGAG 240
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     614 actcagagtgatgaagagaatgggcgtgcctgtgaaatgaatggggaagaatgtgcggag 673

Qy     241 GATTTACGAATGCTTGATGCCTCGGGAGAGAAAATGAATGGCTCCCACAGGGACCAAGGC 300
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     674 gatttacgaatgcttgatgcctcgggagagaaaatgaatggctcccacagggaccaaggc 733

Qy     301 AGCTCGGCTTTGTCGGGAGTTGGAGGCATTCGACTTCCTAACGGAAAACTAAAGTGTGAT 360
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     734 agctcggctttgtcgggagttggaggcattcgacttcctaacggaaaactaaagtgtgat 793

Qy     361 ATCTGTGGGATCATTTGCATCGGGCCCAATGTGCTCATGGTTCACAAAAGAAGCCACACT 420
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     794 atctgtgggatcatttgcatcgggcccaatgtgctcatggttcacaaaagaagccacact 853

Qy     421 GGAGAACGGCCCTTCCAGTGCAATCAGTGCGGGGCCTCATTCACCCAGAAGGGCAACCTG 480
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     854 ggagaacggcccttccagtgcaatcagtgcggggcctcattcacccagaagggcaacctg 913

Qy     481 CTCCGGCACATCAAGCTGCATTCCGGGGAGAAGCCCTTCAAATGCCACCTCTGCAACTAC 540
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     914 ctccggcacatcaagctgcattccggggagaagcccttcaaatgccacctctgcaactac 973

Qy     541 GCCTGCCGCCGGAGGGACGCCCTCACTGGCCACCTGAGGACGCACTCCGTTGGTAAACCT 600
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     974 gcctgccgccggagggacgccctcactggccacctgaggacgcactccgttggtaaacct 1033

Qy     601 CACAAATGTGGATATTGTGGCCGAAGCTATAAACAGCGAACGTCTTTAGAGGAACATAAA 660
           ||||||||||||||||||||||||||||||||||||||  ||||||||||||||||||||
Db    1034 cacaaatgtggatattgtggccgaagctataaacagcgaagctctttagaggaacataaa 1093

Qy     661 GAGCGCTGCCACAACTACTTGGAAAGCATGGGCCTTCCGGGCACACTGTACCCAGTCATT 720
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1094 gagcgctgccacaactacttggaaagcatgggccttccgggcacactgtacccagtcatt 1153

Qy     721 AAAGAAGAAACTAAGCACAGTGAAATGGCAGAAGACCTGTGCAAGATAGGATCAGAGAGA 780
           |||||||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||
Db    1154 aaagaagaaactaatcacagtgaaatggcagaagacctgtgcaagataggatcagagaga 1213

Qy     781 TCTCTCGTGCTGGACAGACTAGCAAGTAATGTCGCCAAACGTAAGAGCTCTATGCCTCAG 840
           |||||||||||||||||||||||||| ||||||||||||||||||||||||||||||||||||
Db    1214 tctctcgtgctggacagactagcaagtaacgtcgccaaacgtaagagctctatgcctcag 1273

Qy     841 AAATTTCTTGGGGACAAGGGCCTGTCCGACACGCCCTACGA---CAGTGCCACGTACGAG 897
           |||||||||||||||||||||||||||||||||||||||||   ||| ||||  ||||||
Db    1274 aaatttcttggggacaagggcctgtccgacacgccctacgacagcagcgccagctacgag 1333

Qy     898 AAGGAGAACGAAATGATGAAGTCCCACGTGATGGACCAAGCCATCAACAACGCCATCAAC 957
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1334 aaggagaacgaaatgatgaagtcccacgtgatggaccaagccatcaacaacgccatcaac 1393

Qy     958 TACCTGGGGGCCGAGTCCCTGCGCCCGCTGGTGCAGACGCCCCCGGGCGGTTCCGAGGTG 1017
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1394 tacctgggggccgagtccctgcgcccgctggtgcagacgccccgggcggttccgaggtg 1453

Qy    1018 GTCCCGGTCATCAGCCCGATGTACCAGCTGCACA---GGCGCTCGGAGGGCACCCCGCGC 1074
           ||||||||||||||||||||||||||||||||    || | ||||||||||||||||||
Db    1454 gtcccggtcatcagcccgatgtaccagctgcacaagccgctcgcggagggcaccccgcgc 1513

Qy    1075 TCCAACCACTCGGCCCAGGACAGCGCCGTGGAGTACCTGCTGCTGCTCTCCAAGGCCAAG 1134
           |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db    1514 tccaaccactcggcccaggacagcgccgtggagaacctgctgctgctctccaaggccaag 1573

Qy    1135 TTGGTGCCCTCGGAGCGCGAGGCGTCCCCGAGCAACAGCTGCCAAGACTCCACGGACACC 1194
           |||||||||||||||||||||||||||||||||||||||||||| ||||||||||||||||||
Db    1574 ttggtgccctcggagcgcgaggcgtccccgagcaacagctgtcaagactccacggacacc 1633

Qy    1195 GAGAGCAACAACGAGGAGCAGCGCAGCGGTCTTATCTACCTGACCAACCACATCGCCCGA 1254
           |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db    1634 gagagcaacaacgaggagcagcgcagcggtctcatctacctgaccaaccacatcgccccg 1693

Qy    1255 CGCGCGC---AACGCGTGTCGCTCAAGGAGGAGCACCGCGCCTACGACCTGCTGCGCGCC 1311
           | |||||   |  || ||||||||||||||||||||||||||||||||||||||||||||
Db    1694 cacgcgcgcaacggcttgtcgctcaaggaggagcaccgcgcctacgacctgctgcgcgcc 1753

Qy    1312 GCCTCCGAGAACTCGCAGGACGCGCTCCGCGTGGTCAGCACCAGCGGGGAGCAGATGAAG 1371
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1754 gcctccgagaactcgcaggacgcgctccgcgtggtcagcaccagcggggagcagatgaag 1813

Qy    1372 GTGTACAAGTGCGAACACTGCCGGGTGCTCTTCCTGGATCACGTCATGTACACCATCCAC 1431
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1814 gtgtacaagtgcgaacactgccgggtgctcttcctggatcacgtcatgtacaccatccac 1873

Qy    1432 ATGGGCTGCCACGGCTTCCGTGATCCTTTTGAGTGCAACATGTGCGGCTACCACAGCCAG 1491
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1874 atgggctgccacggcttccgtgatccttttgagtgcaacatgtgcggctaccacagccag 1933

Qy    1492 GACCGGTACGAGTTCTCGTCGCACATAACGCGAGGGGAGCACCGCTTCCACATGAGCTAA 1551
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1934 gaccggtacgagttctcgtcgcacataacgcgaggggagcaccgcttccacatgagctaa 1993
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RESULT 9
US-09-698-014-4890
; Sequence 4890, Application US/09698014
; GENERAL INFORMATION:
; APPLICANT: Gearing, David P.
; APPLICANT: Holtzman, Douglas A.
; APPLICANT: Kingsbury, Gillian A.
; TITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES
; TITLE OF INVENTION: THEREFOR
; FILE REFERENCE: 1600.2014-001
; CURRENT APPLICATION NUMBER: US/09/698,014
; CURRENT FILING DATE: 2000-10-27
; PRIOR APPLICATION NUMBER: 60/162,363
; PRIOR FILING DATE: 1999-10-29
; NUMBER OF SEQ ID NOS: 6098
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 4890
; LENGTH: 4786
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(4786)
; OTHER INFORMATION: n = A,T,C or G
US-09-698-014-4890

Query Match 94.7%; Score 1468.8; DB 27; Length 4786;
Best Local Similarity 97.7%; Pred. No. 1.8e-297;
Matches 1524; Conservative 0; Mismatches 27; Indels 9; Gaps 3;

```
Qy       1 ATGGATGCTGACGAGGGTCAAGACATGTCTTTCTCATCAGGGAAGGAAAGCCCCCCTGTA 60
           |||||||||| ||||||||||||||||||        |||||||||||||||||||||||
Db     434 atggatgctgatgagggtcaagacatgtcccaagtttcagggaaggaaagccccccctgta 493

Qy      61 AGCGATACTCCAGATGAGGGCGATGAGCCCATGCCGATCCCCGAGGACCTCTCCACCACC 120
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     494 agcgatactccagatgagggcgatgagcccatgccgatccccgaggacctctccaccacc 553

Qy     121 TCGGGAGGACAGCAAAGCTCCAAGAGTGACAGAGTCGTGGCCAGTAATGTTAAAGTAGAG 180
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     554 tcgggaggacagcaaagctccaagagtgacagagtcgtggccagtaatgttaaagtagag 613

Qy     181 ACTCAGAGTGATGAAGAGAATGGGCGTGCCTGTGAAATGAATGGGGAAGAATGTGCGGAG 240
```

```
Db    614 actcagagtgatgaagagaatgggcgtgcctgtgaaatgaatggggaagaatgtgcggag 673

Qy    241 GATTTACGAATGCTTGATGCCTCGGGAGAGAAAATGAATGGCTCCCACAGGGACCAAGGC 300

Db    674 gatttacgaatgcttgatgcctcgggagagaaaatgaatggctcccacagggaccaaggc 733

Qy    301 AGCTCGGCTTTGTCGGGAGTTGGAGGCATTCGACTTCCTAACGGAAAACTAAAGTGTGAT 360

Db    734 agctcggctttgtcgggagttggaggcattcgacttcctaacggaaaactaaagtgtgat 793

Qy    361 ATCTGTGGGATCATTTGCATCGGGCCCAATGTGCTCATGGTTCACAAAAGAAGCCACACT 420

Db    794 atctgtgggatcatttgcatcgggcccaatgtgctcatggttcacaaaagaagccacact 853

Qy    421 GGAGAACGGCCCTTCCAGTGCAATCAGTGCGGGGCCTCATTCACCCAGAAGGGCAACCTG 480

Db    854 ggagaacggcccttccagtgcaatcagtgcggggcctcattcacccagaagggcaacctg 913

Qy    481 CTCCGGCACATCAAGCTGCATTCCGGGGAGAAGCCCTTCAAATGCCACCTCTGCAACTAC 540

Db    914 ctccggcacatcaagctgcattccggggagaagcccttcaaatgccacctctgcaactac 973

Qy    541 GCCTGCCGCCGGAGGGACGCCCTCACTGGCCACCTGAGGACGCACTCCGTTGGTAAACCT 600

Db    974 gcctgccgccggagggacgccctcactggccacctgaggacgcactccgttggtaaacct 1033

Qy    601 CACAAATGTGGATATTGTGGCCGAAGCTATAAACAGCGAACGTCTTTAGAGGAACATAAA 660

Db   1034 cacaaatgtggatattgtggccgaagctataaacagcgaagctctttagaggaacataaa 1093

Qy    661 GAGCGCTGCCACAACTACTTGGAAAGCATGGGCCTTCCGGGCACACTGTACCCAGTCATT 720

Db   1094 gagcgctgccacaactacttggaaagcatgggccttccgggcacactgtacccagtcatt 1153

Qy    721 AAAGAAGAAACTAAGCACAGTGAAATGGCAGAAGACCTGTGCAAGATAGGATCAGAGAGA 780

Db   1154 aaagaagaaactaatcacagtgaaatggcagaagacctgtgcaagataggatcagagaga 1213

Qy    781 TCTCTCGTGCTGGACAGACTAGCAAGTAATGTCGCCAAACGTAAGAGCTCTATGCCTCAG 840

Db   1214 tctctcgtgctggacagactagcaagtaacgtcgccaaacgtaagagctctatgcctcag 1273

Qy    841 AAATTTCTTGGGGACAAGGGCCTGTCCGACACGCCCTACGA---CAGTGCCACGTACGAG 897

Db   1274 aaatttcttggggacaagggcctgtccgacacgccctacgacagcagcgccagctacgag 1333

Qy    898 AAGGAGAACGAAATGATGAAGTCCCACGTGATGGACCAAGCCATCAACAACGCCATCAAC 957

Db   1334 aaggagaacgaaatgatgaagtcccacgtgatggaccaagccatcaacaacgccatcaac 1393

Qy    958 TACCTGGGGGCCGAGTCCCTGCGCCCGCTGGTGCAGACGCCCCCGGGCGGTTCCGAGGTG 1017

Db   1394 tacctgggggccgagtccctgcgcccgctggtgcagacgccccgggcggttccgaggtg 1453

Qy   1018 GTCCCGGTCATCAGCCCGATGTACCAGCTGCACA---GGCGCTCGGAGGGCACCCCGCGC 1074

Db   1454 gtcccggtcatcagcccgatgtaccagctgcacaagccgctcgcggagggcaccccgcgc 1513

Qy   1075 TCCAACCACTCGGCCCAGGACAGCGCCGTGGAGTACCTGCTGCTGCTCTCCAAGGCCAAG 1134

Db   1514 tccaaccactcggcccaggacagcgccgtggagaacctgctgctgctctccaaggccaag 1573

Qy   1135 TTGGTGCCCTCGGAGCGCGAGGCGTCCCCGAGCAACAGCTGCCAAGACTCCACGGACACC 1194

Db   1574 ttggtgccctcggagcgcgaggcgtccccgagcaacagctgtcaagactccacggacacc 1633

Qy   1195 GAGAGCAACAACGAGGAGCAGCGCAGCGGTCTTATCTACCTGACCAACCACATCGCCCGA 1254

Db   1634 gagagcaacaacgaggagcagcgcagcggtctcatctacctgaccaaccacatcgccccg 1693

Qy   1255 CGCGCGC---AACGCGTGTCGCTCAAGGAGGAGCACCGCGCCTACGACCTGCTGCGCGCC 1311

Db   1694 cacgcgcgcaacggcttgtcgctcaaggaggagcaccgcgcctacgacctgctgcgcgcc 1753

Qy   1312 GCCTCCGAGAACTCGCAGGACGCGCTCCGCGTGGTCAGCACCAGCGGGGAGCAGATGAAG 1371

Db   1754 gcctccgagaactcgcaggacgcgctccgcgtggtcagcaccagcggggagcagatgaag 1813

Qy   1372 GTGTACAAGTGCGAACACTGCCGGGTGCTCTTCCTGGATCACGTCATGTACACCATCCAC 1431

Db   1814 gtgtacaagtgcgaacactgccgggtgctcttcctggatcacgtcatgtacaccatccac 1873

Qy   1432 ATGGGCTGCCACGGCTTCCGTGATCCTTTTGAGTGCAACATGTGCGGCTACCACAGCCAG 1491

Db   1874 atgggctgccacggcttccgtgatccttttgagtgcaacatgtgcggctaccacagccag 1933

Qy   1492 GACCGGTACGAGTTCTCGTCGCACATAACGCGAGGGGAGCACCGCTTCCACATGAGCTAA 1551

Db   1934 gaccggtacgagttctcgtcgcacataacgcgaggggagcaccgcttccacatgagctaa 1993
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RESULT 10
US-09-717-350-5922
; Sequence 5922, Application US/09717350
; GENERAL INFORMATION:
; APPLICANT: Gearing, David P.
; APPLICANT: Villeval, Jean-Luc
; APPLICANT: Fraser, Christopher C.
; TITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES
; TITLE OF INVENTION: THEREFOR
; FILE REFERENCE: 1600.2034-001
; CURRENT APPLICATION NUMBER: US/09/717,350
; CURRENT FILING DATE: 2000-11-21
; PRIOR APPLICATION NUMBER: 60/167,885
; PRIOR FILING DATE: 1999-11-24
; NUMBER OF SEQ ID NOS: 5955
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 5922
; LENGTH: 4786
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(4786)
; OTHER INFORMATION: n = A,T,C or G
US-09-717-350-5922

Query Match 94.7%; Score 1468.8; DB 28; Length 4786;
Best Local Similarity 97.7%; Pred. No. 1.8e-297;
Matches 1524; Conservative 0; Mismatches 27; Indels 9; Gaps 3;

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Qy      1 ATGGATGCTGACGAGGGTCAAGACATGTCTTTCTCATCAGGGAAGGAAAGCCCCCCTGTA 60

Db    434 atggatgctgatgagggtcaagacatgtcccaagtttcagggaaggaaagccccccctgta 493

Qy     61 AGCGATACTCCAGATGAGGGCGATGAGCCCATGCCGATCCCCGAGGACCTCTCCACCACC 120

Db    494 agcgatactccagatgagggcgatgagcccatgccgatccccgaggacctctccaccacc 553

Qy    121 TCGGGAGGACAGCAAAGCTCCAAGAGTGACAGAGTCGTGGCCAGTAATGTTAAAGTAGAG 180

Db    554 tcgggaggacagcaaagctccaagagtgacagagtcgtggccagtaatgttaaagtagag 613

Qy    181 ACTCAGAGTGATGAAGAGAATGGGCGTGCCTGTGAAATGAATGGGGAAGAATGTGCGGAG 240

Db    614 actcagagtgatgaagagaatgggcgtgcctgtgaaatgaatggggaagaatgtgcggag 673

Qy    241 GATTTACGAATGCTTGATGCCTCGGGAGAGAAAATGAATGGCTCCCACAGGGACCAAGGC 300

Db    674 gatttacgaatgcttgatgcctcgggagagaaaatgaatggctcccacagggaccaaggc 733

Qy    301 AGCTCGGCTTTGTCGGGAGTTGGAGGCATTCGACTTCCTAACGGAAAACTAAAGTGTGAT 360

Db    734 agctcggctttgtcgggagttggaggcattcgacttcctaacggaaaactaaagtgtgat 793
```

```
Qy     361 ATCTGTGGGATCATTTGCATCGGGCCCAATGTGCTCATGGTTCACAAAAGAAGCCACACT 420
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     794 atctgtgggatcatttgcatcgggcccaatgtgctcatggttcacaaaagaagccacact 853

Qy     421 GGAGAACGGCCCTTCCAGTGCAATCAGTGCGGGGCCTCATTCACCCAGAAGGGCAACCTG 480
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     854 ggagaacggcccttccagtgcaatcagtgcggggcctcattcacccagaagggcaacctg 913

Qy     481 CTCCGGCACATCAAGCTGCATTCCGGGGAGAAGCCCTTCAAATGCCACCTCTGCAACTAC 540
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     914 ctccggcacatcaagctgcattccggggagaagcccttcaaatgccacctctgcaactac 973

Qy     541 GCCTGCCGCCGGAGGGACGCCCTCACTGGCCACCTGAGGACGCACTCCGTTGGTAAACCT 600
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     974 gcctgccgccggagggacgccctcactggccacctgaggacgcactccgttggtaaacct 1033

Qy     601 CACAAATGTGGATATTGTGGCCGAAGCTATAAACAGCGAACGTCTTTAGAGGAACATAAA 660
           ||||||||||||||||||||||||||||||||||||  ||||||||||||||||||||
Db    1034 cacaaatgtggatattgtggccgaagctataaacagcgaagctctttagaggaacataaa 1093

Qy     661 GAGCGCTGCCACAACTACTTGGAAAGCATGGGCCTTCCGGGCACACTGTACCCAGTCATT 720
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1094 gagcgctgccacaactacttggaaagcatgggccttccgggcacactgtacccagtcatt 1153

Qy     721 AAAGAAGAAACTAAGCACAGTGAAATGGCAGAAGACCTGTGCAAGATAGGATCAGAGAGA 780
           |||||||||||||| ||||||||||||||||||||||||||||||||||||||||||||||||
Db    1154 aaagaagaaactaatcacagtgaaatggcagaagacctgtgcaagataggatcagagaga 1213

Qy     781 TCTCTCGTGCTGGACAGACTAGCAAGTAATGTCGCCAAACGTAAGAGCTCTATGCCTCAG 840
           |||||||||||||||||||||||||| ||||||||||||||||||||||||||||||||||||
Db    1214 tctctcgtgctggacagactagcaagtaacgtcgccaaacgtaagagctctatgcctcag 1273

Qy     841 AAATTTCTTGGGGACAAGGGCCTGTCCGACACGCCCTACGA---CAGTGCCACGTACGAG 897
           |||||||||||||||||||||||||||||||||||||||||   ||| ||||  ||||||
Db    1274 aaatttcttggggacaagggcctgtccgacacgccctacgacagcagcgccagctacgag 1333

Qy     898 AAGGAGAACGAAATGATGAAGTCCCACGTGATGGACCAAGCCATCAACAACGCCATCAAC 957
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1334 aaggagaacgaaatgatgaagtcccacgtgatggaccaagccatcaacaacgccatcaac 1393

Qy     958 TACCTGGGGGCCGAGTCCCTGCGCCCGCTGGTGCAGACGCCCCCGGGCGGTTCCGAGGTG 1017
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1394 tacctgggggccgagtccctgcgcccgctggtgcagacgccccgggcggttccgaggtg 1453

Qy    1018 GTCCCGGTCATCAGCCCGATGTACCAGCTGCACA---GGCGCTCGGAGGGCACCCCGCGC 1074
           ||||||||||||||||||||||||||||||||    || | ||||||||||||||||||
Db    1454 gtcccggtcatcagcccgatgtaccagctgcacaagccgctcgcggagggcaccccgcgc 1513

Qy    1075 TCCAACCACTCGGCCCAGGACAGCGCCGTGGAGTACCTGCTGCTGCTCTCCAAGGCCAAG 1134
           |||||||||||||||||||||||||||||||||||| ||||||||||||||||||||||||
Db    1514 tccaaccactcggcccaggacagcgccgtggagaacctgctgctgctctccaaggccaag 1573

Qy    1135 TTGGTGCCCTCGGAGCGCGAGGCGTCCCCGAGCAACAGCTGCCAAGACTCCACGGACACC 1194
           |||||||||||||||||||||||||||||||||||| ||||||||||||||||||||||||
Db    1574 ttggtgccctcggagcgcgaggcgtccccgagcaacagctgtcaagactccacggacacc 1633

Qy    1195 GAGAGCAACAACGAGGAGCAGCGCAGCGGTCTTATCTACCTGACCAACCACATCGCCCGA 1254
           |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db    1634 gagagcaacaacgaggagcagcgcagcggtctcatctacctgaccaaccacatcgccccg 1693

Qy    1255 CGCGCGC---AACGCGTGTCGCTCAAGGAGGAGCACCGCGCCTACGACCTGCTGCGCGCC 1311
           | |||||   |  || ||||||||||||||||||||||||||||||||||||||||||||
Db    1694 cacgcgcgcaacggcttgtcgctcaaggaggagcaccgcgcctacgacctgctgcgcgcc 1753

Qy    1312 GCCTCCGAGAACTCGCAGGACGCGCTCCGCGTGGTCAGCACCAGCGGGGAGCAGATGAAG 1371
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1754 gcctccgagaactcgcaggacgcgctccgcgtggtcagcaccagcggggagcagatgaag 1813

Qy    1372 GTGTACAAGTGCGAACACTGCCGGGTGCTCTTCCTGGATCACGTCATGTACACCATCCAC 1431
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1814 gtgtacaagtgcgaacactgccgggtgctcttcctggatcacgtcatgtacaccatccac 1873

Qy    1432 ATGGGCTGCCACGGCTTCCGTGATCCTTTTGAGTGCAACATGTGCGGCTACCACAGCCAG 1491
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1874 atgggctgccacggcttccgtgatccttttgagtgcaacatgtgcggctaccacagccag 1933

Qy    1492 GACCGGTACGAGTTCTCGTCGCACATAACGCGAGGGGAGCACCGCTTCCACATGAGCTAA 1551
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1934 gaccggtacgagttctcgtcgcacataacgcgaggggagcaccgcttccacatgagctaa 1993
```

RESULT 11
US-09-726-802-2123
; Sequence 2123, Application US/09726802
; GENERAL INFORMATION:
; APPLICANT: Acton, Susan L.
; APPLICANT: Falb, Dean A.
; APPLICANT: .
; TITLE OF INVENTION: NOVEL NUCLEIC ACID MOLECULES AND USES
; TITLE OF INVENTION: THEREFOR
; FILE REFERENCE: 1600.2025-001
; CURRENT APPLICATION NUMBER: US/09/726,802
; CURRENT FILING DATE: 2000-11-30
; PRIOR APPLICATION NUMBER: 60/168,012
; PRIOR FILING DATE: 1999-11-30
; NUMBER OF SEQ ID NOS: 2872
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 2123
; LENGTH: 4786
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)...(4786)
; OTHER INFORMATION: n = A,T,C or G
US-09-726-802-2123

Query Match 94.7%; Score 1468.8; DB 29; Length 4786;
Best Local Similarity 97.7%; Pred. No. 1.8e-297;
Matches 1524; Conservative 0; Mismatches 27; Indels 9; Gaps 3;

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Qy       1 ATGGATGCTGACGAGGGTCAAGACATGTCTTTCTCATCAGGGAAGGAAAGCCCCCCTGTA 60
           |||||||||| |||||||||||||||||       |||||||||||||||||||||||||
Db     434 atggatgctgatgagggtcaagacatgtcccaagtttcagggaaggaaagccccccctgta 493

Qy      61 AGCGATACTCCAGATGAGGGCGATGAGCCCATGCCGATCCCCGAGGACCTCTCCACCACC 120
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     494 agcgatactccagatgagggcgatgagcccatgccgatccccgaggacctctccaccacc 553

Qy     121 TCGGGAGGACAGCAAAGCTCCAAGAGTGACAGAGTCGTGGCCAGTAATGTTAAAGTAGAG 180
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     554 tcgggaggacagcaaagctccaagagtgacagagtcgtggccagtaatgttaaagtagag 613

Qy     181 ACTCAGAGTGATGAAGAGAATGGGCGTGCCTGTGAAATGAATGGGGAAGAATGTGCGGAG 240
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     614 actcagagtgatgaagagaatgggcgtgcctgtgaaatgaatggggaagaatgtgcggag 673

Qy     241 GATTTACGAATGCTTGATGCCTCGGGAGAGAAAATGAATGGCTCCCACAGGGACCAAGGC 300
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     674 gatttacgaatgcttgatgcctcgggagagaaaatgaatggctcccacagggaccaaggc 733

Qy     301 AGCTCGGCTTTGTCGGGAGTTGGAGGCATTCGACTTCCTAACGGAAAACTAAAGTGTGAT 360
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     734 agctcggctttgtcgggagttggaggcattcgacttcctaacggaaaactaaagtgtgat 793

Qy     361 ATCTGTGGGATCATTTGCATCGGGCCCAATGTGCTCATGGTTCACAAAAGAAGCCACACT 420
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     794 atctgtgggatcatttgcatcgggcccaatgtgctcatggttcacaaaagaagccacact 853

Qy     421 GGAGAACGGCCCTTCCAGTGCAATCAGTGCGGGGCCTCATTCACCCAGAAGGGCAACCTG 480
           ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     854 ggagaacggcccttccagtgcaatcagtgcggggcctcattcacccagaagggcaacctg 913

Qy     481 CTCCGGCACATCAAGCTGCATTCCGGGGAGAAGCCCTTCAAATGCCACCTCTGCAACTAC 540
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```
Db    914 ctccggcacatcaagctgcattccggggagaagcccttcaaatgccacctctgcaactac 973

Qy    541 GCCTGCCGCCGGAGGGACGCCCTCACTGGCCACCTGAGGACGCACTCCGTTGGTAAACCT 600

Db    974 gcctgccgccggagggacgccctcactggccacctgaggacgcactccgttggtaaacct 1033

Qy    601 CACAAATGTGGATATTGTGGCCGAAGCTATAAACAGCGAACGTCTTTAGAGGAACATAAA 660

Db   1034 cacaaatgtggatattgtggccgaagctataaacagcgaagctctttagaggaacataaa 1093

Qy    661 GAGCGCTGCCACAACTACTTGGAAAGCATGGGCCTTCCGGGCACACTGTACCCAGTCATT 720

Db   1094 gagcgctgccacaactacttggaaagcatgggccttccgggcacactgtacccagtcatt 1153

Qy    721 AAAGAAGAAACTAAGCACAGTGAAATGGCAGAAGACCTGTGCAAGATAGGATCAGAGAGA 780

Db   1154 aaagaagaaactaatcacagtgaaatggcagaagacctgtgcaagataggatcagagaga 1213

Qy    781 TCTCTCGTGCTGGACAGACTAGCAAGTAATGTCGCCAAACGTAAGAGCTCTATGCCTCAG 840

Db   1214 tctctcgtgctggacagactagcaagtaacgtcgccaaacgtaagagctctatgcctcag 1273

Qy    841 AAATTTCTTGGGGACAAGGGCCTGTCCGACACGCCCTACGA---CAGTGCCACGTACGAG 897

Db   1274 aaatttcttggggacaagggcctgtccgacacgccctacgacagcagcgccagctacgag 1333

Qy    898 AAGGAGAACGAAATGATGAAGTCCCACGTGATGGACCAAGCCATCAACAACGCCATCAAC 957

Db   1334 aaggagaacgaaatgatgaagtcccacgtgatggaccaagccatcaacaacgccatcaac 1393

Qy    958 TACCTGGGGGCCGAGTCCCTGCGCCCGCTGGTGCAGACGCCCCCGGGCGGTTCCGAGGTG 1017

Db   1394 tacctgggggccgagtccctgcgcccgctggtgcagacgccccgggcggttccgaggtg 1453

Qy   1018 GTCCCGGTCATCAGCCCGATGTACCAGCTGCACA---GGCGCTCGGAGGGCACCCCGCGC 1074

Db   1454 gtcccggtcatcagcccgatgtaccagctgcacaagccgctcgcggagggcaccccgcgc 1513

Qy   1075 TCCAACCACTCGGCCCAGGACAGCGCCGTGGAGTACCTGCTGCTGCTCTCCAAGGCCAAG 1134

Db   1514 tccaaccactcggcccaggacagcgccgtggagaacctgctgctgctctccaaggccaag 1573

Qy   1135 TTGGTGCCCTCGGAGCGCGAGGCGTCCCCGAGCAACAGCTGCCAAGACTCCACGGACACC 1194

Db   1574 ttggtgccctcggagcgcgaggcgtccccgagcaacagctgtcaagactccacggacacc 1633

Qy   1195 GAGAGCAACAACGAGGAGCAGCGCAGCGGTCTTATCTACCTGACCAACCACATCGCCCGA 1254

Db   1634 gagagcaacaacgaggagcagcgcagcggtctcatctacctgaccaaccacatcgccccg 1693

Qy   1255 CGCGCGC---AACGCGTGTCGCTCAAGGAGGAGCACCGCGCCTACGACCTGCTGCGCGCC 1311

Db   1694 cacgcgcgcaacggcttgtcgctcaaggaggagcaccgcgcctacgacctgctgcgcgcc 1753

Qy   1312 GCCTCCGAGAACTCGCAGGACGCGCTCCGCGTGGTCAGCACCAGCGGGGAGCAGATGAAG 1371

Db   1754 gcctccgagaactcgcaggacgcgctccgcgtggtcagcaccagcggggagcagatgaag 1813

Qy   1372 GTGTACAAGTGCGAACACTGCCGGGTGCTCTTCCTGGATCACGTCATGTACACCATCCAC 1431

Db   1814 gtgtacaagtgcgaacactgccgggtgctcttcctggatcacgtcatgtacaccatccac 1873

Qy   1432 ATGGGCTGCCACGGCTTCCGTGATCCTTTTGAGTGCAACATGTGCGGCTACCACAGCCAG 1491

Db   1874 atgggctgccacggcttccgtgatccttttgagtgcaacatgtgcggctaccacagccag 1933

Qy   1492 GACCGGTACGAGTTCTCGTCGCACATAACGCGAGGGGAGCACCGCTTCCACATGAGCTAA 1551

Db   1934 gaccggtacgagttctcgtcgcacataacgcgaggggagcaccgcttccacatgagctaa 1993

RESULT 12
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US-09-726-811-4096
; Sequence 4096, Application US/09726811
; GENERAL INFORMATION:
;  APPLICANT:  Gutierrez-Ramos, Jose-Carlos
;  APPLICANT:  Weich, Nadine S.
;  APPLICANT:  Wen, Danyi
;  TITLE OF INVENTION:  NOVEL NUCLEIC ACID MOLECULES AND USES
;  TITLE OF INVENTION:  THEREFOR
;  FILE REFERENCE:  1600.2027-001
;  CURRENT APPLICATION NUMBER: US/09/726,811
;  CURRENT FILING DATE:  2000-11-30
;  PRIOR APPLICATION NUMBER:  60/168,136
;  PRIOR FILING DATE:  1999-11-30
;  NUMBER OF SEQ ID NOS:  5515
;  SOFTWARE:  FastSEQ for Windows Version 4.0
; SEQ ID NO 4096
;   LENGTH: 4786
;   TYPE: DNA
;   ORGANISM: Homo sapiens
;   FEATURE:
;   NAME/KEY: misc_feature
;   LOCATION: (1)...(4786)
;   OTHER INFORMATION: n = A,T,C or G
US-09-726-811-4096
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Query Match 94.7%; Score 1468.8; DB 29; Length 4786;
Best Local Similarity 97.7%; Pred. No. 1.8e-297;
Matches 1524; Conservative 0; Mismatches 27; Indels 9; Gaps 3;

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Qy      1 ATGGATGCTGACGAGGGTCAAGACATGTCTTTCTCATCAGGGAAGGAAAGCCCCCCTGTA 60

Db    434 atggatgctgatgagggtcaagacatgtcccaagtttcagggaaggaaagccccctgta 493

Qy     61 AGCGATACTCCAGATGAGGGCGATGAGCCCATGCCGATCCCCGAGGACCTCTCCACCACC 120

Db    494 agcgatactccagatgagggcgatgagcccatgccgatccccgaggacctctccaccacc 553

Qy    121 TCGGGAGGACAGCAAAGCTCCAAGAGTGACAGAGTCGTGGCCAGTAATGTTAAAGTAGAG 180

Db    554 tcgggaggacagcaaagctccaagagtgacagagtcgtggccagtaatgttaaagtagag 613

Qy    181 ACTCAGAGTGATGAAGAGAATGGGCGTGCCTGTGAAATGAATGGGGAAGAATGTGCGGAG 240

Db    614 actcagagtgatgaagagaatgggcgtgcctgtgaaatgaatggggaagaatgtgcggag 673

Qy    241 GATTTACGAATGCTTGATGCCTCGGGAGAGAAAATGAATGGCTCCCACAGGGACCAAGGC 300

Db    674 gatttacgaatgcttgatgcctcgggagagaaaatgaatggctcccacagggaccaaggc 733

Qy    301 AGCTCGGCTTTGTCGGGAGTTGGAGGCATTCGACTTCCTAACGGAAAACTAAAGTGTGAT 360

Db    734 agctcggctttgtcgggagttggaggcattcgacttcctaacggaaaactaaagtgtgat 793

Qy    361 ATCTGTGGGATCATTTGCATCGGGCCCAATGTGCTCATGGTTCACAAAAGAAGCCACACT 420

Db    794 atctgtgggatcatttgcatcgggcccaatgtgctcatggttcacaaaagaagccacact 853

Qy    421 GGAGAACGGCCCTTCCAGTGCAATCAGTGCGGGGCCTCATTCACCCAGAAGGGCAACCTG 480

Db    854 ggagaacggcccttccagtgcaatcagtgcggggcctcattcacccagaagggcaacctg 913

Qy    481 CTCCGGCACATCAAGCTGCATTCCGGGGAGAAGCCCTTCAAATGCCACCTCTGCAACTAC 540

Db    914 ctccggcacatcaagctgcattccggggagaagcccttcaaatgccacctctgcaactac 973

Qy    541 GCCTGCCGCCGGAGGGACGCCCTCACTGGCCACCTGAGGACGCACTCCGTTGGTAAACCT 600

Db    974 gcctgccgccggagggacgccctcactggccacctgaggacgcactccgttggtaaacct 1033

Qy    601 CACAAATGTGGATATTGTGGCCGAAGCTATAAACAGCGAACGTCTTTAGAGGAACATAAA 660

Db   1034 cacaaatgtggatattgtggccgaagctataaacagcgaagctctttagaggaacataaa 1093
```

```
Qy    661 GAGCGCTGCCACAACTACTTGGAAAGCATGGGCCTTCCGGGCACACTGTACCCAGTCATT 720
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1094 gagcgctgccacaactacttggaaagcatgggccttccgggcacactgtacccagtcattt 1153

Qy    721 AAAGAAGAAACTAAGCACAGTGAAATGGCAGAAGACCTGTGCAAGATAGGATCAGAGAGA 780
          ||||||||||||| ||||||||||||||||||||||||||||||||||||||||||||||
Db   1154 aaagaagaaactaatcacagtgaaatggcagaagacctgtgcaagataggatcagagaga 1213

Qy    781 TCTCTCGTGCTGGACAGACTAGCAAGTAATGTCGCCAAACGTAAGAGCTCTATGCCTCAG 840
          |||||||||||||||||||||||||||| ||||||||||||||||||||||||||||||
Db   1214 tctctcgtgctggacagactagcaagtaacgtcgccaaacgtaagagctctatgcctcag 1273

Qy    841 AAATTTCTTGGGGACAAGGGCCTGTCCGACACGCCCTACGA---CAGTGCCACGTACGAG 897
          |||||||||||||||||||||||||||||||||||||||   ||| ||||  ||||||
Db   1274 aaatttcttggggacaagggcctgtccgacacgccctacgacagcagcgccagctacgag 1333

Qy    898 AAGGAGAACGAAATGATGAAGTCCCACGTGATGGACCAAGCCATCAACAACGCCATCAAC 957
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1334 aaggagaacgaaatgatgaagtcccacgtgatggaccaagccatcaacaacgccatcaac 1393

Qy    958 TACCTGGGGGCCGAGTCCCTGCGCCCGCTGGTGCAGACGCCCCCGGGCGGTTCCGAGGTG 1017
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1394 tacctgggggccgagtccctgcgcccgctggtgcagacgccccgggcggttccgaggtg 1453

Qy   1018 GTCCCGGTCATCAGCCCGATGTACCAGCTGCACA---GGCGCTCGGAGGGCACCCCGCGC 1074
          |||||||||||||||||||||||||||||||||   || | ||||||||||||||||||
Db   1454 gtcccggtcatcagcccgatgtaccagctgcacaagccgctcgcggagggcaccccgcgc 1513

Qy   1075 TCCAACCACTCGGCCCAGGACAGCGCCGTGGAGTACCTGCTGCTGCTCTCCAAGGCCAAG 1134
          |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db   1514 tccaaccactcggcccaggacagcgccgtggagaacctgctgctgctctccaaggccaag 1573

Qy   1135 TTGGTGCCCTCGGAGCGCGAGGCGTCCCCGAGCAACAGCTGCCAAGACTCCACGGACACC 1194
          |||||||||||||||||||||||||||||||||||||||| ||||||||||||||||||
Db   1574 ttggtgccctcggagcgcgaggcgtccccgagcaacagctgtcaagactccacggacacc 1633

Qy   1195 GAGAGCAACAACGAGGAGCAGCGCAGCGGTCTTATCTACCTGACCAACCACATCGCCCGA 1254
          |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||
Db   1634 gagagcaacaacgaggagcagcgcagcggtctcatctacctgaccaaccacatcgccccg 1693

Qy   1255 CGCGCGC---AACGCGTGTCGCTCAAGGAGGAGCACCGCGCCTACGACCTGCTGCGCGCC 1311
          | |||||   |  || ||||||||||||||||||||||||||||||||||||||||||||
Db   1694 cacgcgcgcaacggcttgtcgctcaaggaggagcaccgcgcctacgacctgctgcgcgcc 1753

Qy   1312 GCCTCCGAGAACTCGCAGGACGCGCTCCGCGTGGTCAGCACCAGCGGGGAGCAGATGAAG 1371
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1754 gcctccgagaactcgcaggacgcgctccgcgtggtcagcaccagcggggagcagatgaag 1813

Qy   1372 GTGTACAAGTGCGAACACTGCCGGGTGCTCTTCCTGGATCACGTCATGTACACCATCCAC 1431
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1814 gtgtacaagtgcgaacactgccgggtgctcttcctggatcacgtcatgtacaccatccac 1873

Qy   1432 ATGGGCTGCCACGGCTTCCGTGATCCTTTTGAGTGCAACATGTGCGGCTACCACAGCCAG 1491
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1874 atgggctgccacggcttccgtgatcctttgagtgcaacatgtgcggctaccacagccag 1933

Qy   1492 GACCGGTACGAGTTCTCGTCGCACATAACGCGAGGGGAGCACCGCTTCCACATGAGCTAA 1551
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1934 gaccggtacgagttctcgtcgcacataacgcgaggggagcaccgcttccacatgagctaa 1993
```

RESULT 13
US-60-324-185-24401
; Sequence 24401, Application US/60324185
; GENERAL INFORMATION:
; APPLICANT: Morris, MacDonald
; APPLICANT: Lal, Preeti
; APPLICANT: Diep, Dinh
; TITLE OF INVENTION: METHOD FOR THE IDENTIFICATION OF SEQUENCE POLYMORPHISMS USING
; TITLE OF INVENTION: POLYNUCLEOTIDE SEQUENCE DATABASES, AND SINGLE NUCLEOTIDE
; TITLE OF INVENTION: POLYMORPHISMS IDENTIFIED THEREBY
; FILE REFERENCE: GX-0019-1 P
; CURRENT APPLICATION NUMBER: US/60/324,185
; CURRENT FILING DATE: 2001-09-21
; NUMBER OF SEQ ID NOS: 35862
; SOFTWARE: PERL Program
; SEQ ID NO 24401
; LENGTH: 6203
; TYPE: DNA
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; OTHER INFORMATION: Incyte ID No: 345320.3
US-60-324-185-24401

Query Match 94.0%; Score 1457.8; DB 71; Length 6203;
Best Local Similarity 97.6%; Pred. No. 3.8e-295;
Matches 1524; Conservative 0; Mismatches 27; Indels 10; Gaps 4;

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Qy      1 ATGGATGCTGACGAGGGTCAAGACATGTCTTTCTCATCAGGGAAGGAAAGCCCCCCTGTA 60
          |||||||||| ||||||||||||||||||        ||||||||||||||||||||||||
Db    169 atggatgctgatgagggtcaagacatgtcccaagtttcagggaaggaaagccccctgta 228

Qy     61 AGCGATACTCCAGATGAGGGCGATGAGCCCATGCCGATCCCCGAGGACCTCTCCACCACC 120
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    229 agcgatactccagatgagggcgatgagcccatgccgatccccgaggacctctccaccacc 288

Qy    121 TCGGGAGGACAGCAAAGCTCCAAGAGTGACAGAGTCGTGGCCAGTAATGTTAAAGTAGAG 180
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    289 tcgggaggacagcaaagctccaagagtgacagagtcgtggccagtaatgttaaagtagag 348

Qy    181 ACTCAGAGTGATGAAGAGAATGGGCGTGCCTGTGAAATGAATGGGGAAGAATGTGCGGAG 240
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    349 actcagagtgatgaagagaatgggcgtgcctgtgaaatgaatggggaagaatgtgcggag 408

Qy    241 GATTTACGAATGCTTGATGCCTCGGGAGAGAAAATGAATGGCTCCCACAGGGACCAAGGC 300
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    409 gatttacgaatgcttgatgcctcgggagagaaaatgaatggctcccacagggaccaaggc 468

Qy    301 AGCTCGGCTTTGTCGGGAGTTGGAGGCATTCGACTTCCTAACGGA-AAACTAAAGTGTGA 359
          |||||||||||||||||||||||||||||||||||||||||||| ||||||||||||||
Db    469 agctcggctttgtcgggagttggaggcattcgacttcctaacggagaaactaaagtgtga 528

Qy    360 TATCTGTGGGATCATTTGCATCGGGCCCAATGTGCTCATGGTTCACAAAAGAAGCCACAC 419
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    529 tatctgtgggatcatttgcatcgggcccaatgtgctcatggttcacaaaagaagccacac 588

Qy    420 TGGAGAACGGCCCTTCCAGTGCAATCAGTGCGGGGCCTCATTCACCCAGAAGGGCAACCT 479
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    589 tggagaacggcccttccagtgcaatcagtgcggggcctcattcacccagaagggcaacct 648

Qy    480 GCTCCGGCACATCAAGCTGCATTCCGGGGAGAAGCCCTTCAAATGCCACCTCTGCAACTA 539
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    649 gctccggcacatcaagctgcattccggggagaagcccttcaaatgccacctctgcaacta 708

Qy    540 CGCCTGCCGCCGGAGGGACGCCCTCACTGGCCACCTGAGGACGCACTCCGTTGGTAAACC 599
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    709 cgcctgccgccggagggacgccctcactggccacctgaggacgcactccgttggtaaacc 768

Qy    600 TCACAAATGTGGATATTGTGGCCGAAGCTATAAACAGCGAACGTCTTTAGAGGAACATAA 659
          |||||||||||||||||||||||||||||||||||||||||  ||||||||||||||||
Db    769 tcacaaatgtggatattgtggccgaagctataaacagcgaagctctttagaggaacataa 828

Qy    660 AGAGCGCTGCCACAACTACTTGGAAAGCATGGGCCTTCCGGGCACACTGTACCCAGTCAT 719
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    829 agagcgctgccacaactacttggaaagcatgggccttccgggcacactgtacccagtcat 888

Qy    720 TAAAGAAGAAACTAAGCACAGTGAAATGGCAGAAGACCTGTGCAAGATAGGATCAGAGAG 779
          |||||||||||||| ||||||||||||||||||||||||||||||||||||||||||||
Db    889 taaagaagaaactaatcacagtgaaatggcagaagacctgtgcaagataggatcagagag 948

Qy    780 ATCTCTCGTGCTGGACAGACTAGCAAGTAATGTCGCCAAACGTAAGAGCTCTATGCCTCA 839
          |||||||||||||||||||||||||||||| ||||||||||||||||||||||||||||
Db    949 atctctcgtgctggacagactagcaagtaacgtcgccaaacgtaagagctctatgcctca 1008
```

```
Qy    840 GAAATTTCTTGGGGACAAGGGCCTGTCCGACACGCCCTACGA---CAGTGCCACGTACGA 896
          ||||||||||||||||||||||||||||||||||||||||||   ||| ||||  ||||| 
Db   1009 gaaatttcttggggacaagggcctgtccgacacgccctacgacagcagcgccagctacga 1068

Qy    897 GAAGGAGAACGAAATGATGAAGTCCCACGTGATGGACCAAGCCATCAACAACGCCATCAA 956
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db   1069 gaaggagaacgaaatgatgaagtcccacgtgatggaccaagccatcaacaacgccatcaa 1128

Qy    957 CTACCTGGGGGCCGAGTCCCTGCGCCCGCTGGTGCAGACGCCCCCGGGCGGTTCCGAGGT 1016
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db   1129 ctacctgggggccgagtccctgcgcccgctggtgcagacgccccgggcggttccgaggt 1188

Qy   1017 GGTCCCGGTCATCAGCCCGATGTACCAGCTGCACA---GGCGCTCGGAGGGCACCCCGCG 1073
          ||||||||||||||||||||||||||||||||    || | ||||||||||||||||  
Db   1189 ggtcccggtcatcagcccgatgtaccagctgcacaagccgctcgcggagggcaccccgcg 1248

Qy   1074 CTCCAACCACTCGGCCCAGGACAGCGCCGTGGAGTACCTGCTGCTGCTCTCCAAGGCCAA 1133
          |||||||||||||||||||||||||||||||| ||||||||||||||||||||||||||| 
Db   1249 ctccaaccactcggcccaggacagcgccgtggagaacctgctgctgctctccaaggccaa 1308

Qy   1134 GTTGGTGCCCTCGGAGCGCGAGGCGTCCCCGAGCAACAGCTGCCAAGACTCCACGGACAC 1193
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db   1309 gttggtgccctcggagcgcgaggcgtccccgagcaacagctgccaagactccacggacac 1368

Qy   1194 CGAGAGCAACAACGAGGAGCAGCGCAGCGGTCTTATCTACCTGACCAACCACATCGCCCG 1253
          |||||||||||||||||||||||||||||||| |||||||||||||||||||||||||  
Db   1369 cgagagcaacaacgaggagcagcgcagcggtctcatctacctgaccaaccacatcgcccc 1428

Qy   1254 ACGCGCGC---AACGCGTGTCGCTCAAGGAGGAGCACCGCGCCTACGACCTGCTGCGCGC 1310
           | |||||   | | ||||||||||||||||||||||||||||||||||||||||||| 
Db   1429 gcacgcgcgcaacgggctgtcgctcaaggaggagcaccgcgcctacgacctgctgcgcgc 1488

Qy   1311 CGCCTCCGAGAACTCGCAGGACGCGCTCCGCGTGGTCAGCACCAGCGGGGAGCAGATGAA 1370
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db   1489 cgcctccgagaactcgcaggacgcgctccgcgtggtcagcaccagcggggagcagatgaa 1548

Qy   1371 GGTGTACAAGTGCGAACACTGCCGGGTGCTCTTCCTGGATCACGTCATGTACACCATCCA 1430
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db   1549 ggtgtacaagtgcgaacactgccgggtgctcttcctggatcacgtcatgtacaccatcca 1608

Qy   1431 CATGGGCTGCCACGGCTTCCGTGATCCTTTTGAGTGCAACATGTGCGGCTACCACAGCCA 1490
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db   1609 catgggctgccacggcttccgtgatccttttgagtgcaacatgtgcggctaccacagcca 1668

Qy   1491 GGACCGGTACGAGTTCTCGTCGCACATAACGCGAGGGGAGCACCGCTTCCACATGAGCTA 1550
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db   1669 ggaccggtacgagttctcgtcgcacataacgcgaggggagcaccgcttccacatgagcta 1728

Qy   1551 A 1551
          |
Db   1729 a 1729
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RESULT  14
PCT-US99-02559-16
; Sequence 16, Application PC/TUS9902559
; GENERAL INFORMATION:
;  APPLICANT: Shiseido Co., Ltd.
;  TITLE OF INVENTION: AIOLOS GENE
;  FILE REFERENCE: 10287/031WO1
;  CURRENT APPLICATION NUMBER: PCT/US99/02559
;  CURRENT FILING DATE: 1999-02-05
;  EARLIER APPLICATION NUMBER: US 09/019,348
;  EARLIER FILING DATE: 1998-02-05
;  NUMBER OF SEQ ID NOS: 29
;  SOFTWARE: FastSEQ for Windows Version 3.0
; SEQ ID NO 16
;  LENGTH: 1386
;  TYPE: DNA
;  ORGANISM: Mus musculus
;  FEATURE:
;  NAME/KEY: CDS
;  LOCATION: (1)...(1383)
PCT-US99-02559-16
```

Query Match 89.4%; Score 1386; DB 1; Length 1386;
Best Local Similarity 100.0%; Pred. No. 3.5e-280;
Matches 1386; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```
Qy    166 AATGTTAAAGTAGAGACTCAGAGTGATGAAGAGAATGGGCGTGCCTGTGAAATGAATGGG 225
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db      1 aatgttaaagtagagactcagagtgatgaagagaatgggcgtgcctgtgaaatgaatggg 60

Qy    226 GAAGAATGTGCGGAGGATTTACGAATGCTTGATGCCTCGGGAGAGAAAATGAATGGCTCC 285
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db     61 gaagaatgtgcggaggatttacgaatgcttgatgcctcgggagagaaaatgaatggctcc 120

Qy    286 CACAGGGACCAAGGCAGCTCGGCTTTGTCGGGAGTTGGAGGCATTCGACTTCCTAACGGA 345
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db    121 cacagggaccaaggcagctcggctttgtcgggagttggaggcattcgacttcctaacgga 180

Qy    346 AAACTAAAGTGTGATATCTGTGGGATCATTTGCATCGGGCCCAATGTGCTCATGGTTCAC 405
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db    181 aaactaaagtgtgatatctgtgggatcatttgcatcgggcccaatgtgctcatggttcac 240

Qy    406 AAAAGAAGCCACACTGGAGAACGGCCCTTCCAGTGCAATCAGTGCGGGGCCTCATTCACC 465
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db    241 aaaagaagccacactggagaacggcccttccagtgcaatcagtgcggggcctcattcacc 300

Qy    466 CAGAAGGGCAACCTGCTCCGGCACATCAAGCTGCATTCCGGGGAGAAGCCCTTCAAATGC 525
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db    301 cagaagggcaacctgctccggcacatcaagctgcattccggggagaagcccttcaaatgc 360

Qy    526 CACCTCTGCAACTACGCCTGCCGCCGGAGGGACGCCCTCACTGGCCACCTGAGGACGCAC 585
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db    361 cacctctgcaactacgcctgccgccggagggacgccctcactggccacctgaggacgcac 420

Qy    586 TCCGTTGGTAAACCTCACAAATGTGGATATTGTGGCCGAAGCTATAAACAGCGAACGTCT 645
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db    421 tccgttggtaaacctcacaaatgtggatattgtggccgaagctataaacagcgaacgtct 480

Qy    646 TTAGAGGAACATAAAGAGCGCTGCCACAACTACTTGGAAAGCATGGGCCTTCCGGGCACA 705
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db    481 ttagaggaacataaagagcgctgccacaactacttggaaagcatgggccttccgggcaca 540

Qy    706 CTGTACCCAGTCATTAAAGAAGAAACTAAGCACAGTGAAATGGCAGAAGACCTGTGCAAG 765
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db    541 ctgtacccagtcattaaagaagaaactaagcacagtgaaatggcagaagacctgtgcaag 600

Qy    766 ATAGGATCAGAGAGATCTCTCGTGCTGGACAGACTAGCAAGTAATGTCGCCAAACGTAAG 825
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db    601 ataggatcagagagatctctcgtgctggacagactagcaagtaatgtcgccaaacgtaag 660

Qy    826 AGCTCTATGCCTCAGAAATTTCTTGGGGACAAGGGCCTGTCCGACACGCCCTACGACAGT 885
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db    661 agctctatgcctcagaaatttcttggggacaagggcctgtccgacacgccctacgacagt 720

Qy    886 GCCACGTACGAGAAGGAGAACGAAATGATGAAGTCCCACGTGATGGACCAAGCCATCAAC 945
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db    721 gccacgtacgagaaggagaacgaaatgatgaagtcccacgtgatggaccaagccatcaac 780

Qy    946 AACGCCATCAACTACCTGGGGGCCGAGTCCCTGCGCCCGCTGGTGCAGACGCCCCCGGGC 1005
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db    781 aacgccatcaactacctgggggccgagtccctgcgcccgctggtgcagacgccccgggc 840

Qy   1006 GGTTCCGAGGTGGTCCCGGTCATCAGCCCGATGTACCAGCTGCACAGGCGCTCGGAGGGC 1065
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db    841 ggttccgaggtggtcccggtcatcagcccgatgtaccagctgcacaggcgctcggagggc 900

Qy   1066 ACCCCGCGCTCCAACCACTCGGCCCAGGACAGCGCCGTGGAGTACCTGCTGCTGCTCTCC 1125
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||| 
Db    901 accccgcgctccaaccactcggcccaggacagcgccgtggagtacctgctgctgctctcc 960

Qy   1126 AAGGCCAAGTTGGTGCCCTCGGAGCGCGAGGCGTCCCCGAGCAACAGCTGCCAAGACTCC 1185
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          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    961 aaggccaagttggtgccctcggagcgcgaggcgtccccgagcaacagctgccaagactcc 1020

Qy   1186 ACGGACACCGAGAGCAACAACGAGGAGCAGCGCAGCGGTCTTATCTACCTGACCAACCAC 1245
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1021 acggacaccgagagcaacaacgaggagcagcgcagcggtcttatctacctgaccaaccac 1080

Qy   1246 ATCGCCCGACGCGCGCAACGCGTGTCGCTCAAGGAGGAGCACCGCGCCTACGACCTGCTG 1305
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1081 atcgcccgacgcgcgcaacgcgtgtcgctcaaggaggagcaccgcgcctacgacctgctg 1140

Qy   1306 CGCGCCGCCTCCGAGAACTCGCAGGACGCGCTCCGCGTGGTCAGCACCAGCGGGGAGCAG 1365
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1141 cgcgccgcctccgagaactcgcaggacgcgctccgcgtggtcagcaccagcggggagcag 1200

Qy   1366 ATGAAGGTGTACAAGTGCGAACACTGCCGGGTGCTCTTCCTGGATCACGTCATGTACACC 1425
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1201 atgaaggtgtacaagtgcgaacactgccgggtgctcttcctggatcacgtcatgtacacc 1260

Qy   1426 ATCCACATGGGCTGCCACGGCTTCCGTGATCCTTTTGAGTGCAACATGTGCGGCTACCAC 1485
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1261 atccacatgggctgccacggcttccgtgatccttttgagtgcaacatgtgcggctaccac 1320

Qy   1486 AGCCAGGACCGGTACGAGTTCTCGTCGCACATAACGCGAGGGGAGCACCGCTTCCACATG 1545
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db   1321 agccaggaccggtacgagttctcgtcgcacataacgcgaggggagcaccgcttccacatg 1380

Qy   1546 AGCTAA 1551
          ||||||
Db   1381 agctaa 1386


RESULT  15
US-08-238-212A-3
; Sequence 3, Application US/08238212A
;  GENERAL INFORMATION:
;    APPLICANT:  Georgopoulos, Katia A.
;    TITLE OF INVENTION:  IKAROS: A T CELL PATHWAY REGULATORY GENE
;    NUMBER OF SEQUENCES:  164
;    CORRESPONDENCE ADDRESS:
;      ADDRESSEE:  LAHIVE & COCKFIELD
;      STREET:  60 STATE STREET, Suite 510
;      CITY:  BOSTON
;      STATE:  MASSACHUCETTS
;      COUNTRY:  USA
;      ZIP:  02109
;    COMPUTER READABLE FORM:
;      MEDIUM TYPE:  Floppy disk
;      COMPUTER:  IBM PC compatible
;      OPERATING SYSTEM:  PC-DOS/MS-DOS
;      SOFTWARE:  AscII (text)
;    CURRENT APPLICATION DATA:
;      APPLICATION NUMBER:  US/08/238,212A
;      FILING DATE:
;      CLASSIFICATION:  424
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 121,438
;      FILING DATE:  14-SEP-1993
;    PRIOR APPLICATION DATA:
;      APPLICATION NUMBER:  US 946,233
;      FILING DATE:  14-SEP-1992
;    ATTORNEY/AGENT INFORMATION:
;      NAME:  Myers, Paul L.
;      REGISTRATION NUMBER:  35,695
;      REFERENCE/DOCKET NUMBER:  MPG-006CP2
;    TELECOMMUNICATION INFORMATION:
;      TELEPHONE:  (617)227-7400
;      TELEFAX:  (617)227-5941
;  INFORMATION FOR SEQ ID NO:  3:
;    SEQUENCE CHARACTERISTICS:
;      LENGTH:  1386 base pairs
;      TYPE:  nucleic acid
;      STRANDEDNESS:  double
;      TOPOLOGY:  linear
;    MOLECULE TYPE:  cDNA
;    FEATURE:
;      NAME/KEY:  CDS
;      LOCATION:  1..1386
US-08-238-212A-3


  Query Match             89.4%;  Score 1386;  DB 6;  Length 1386;
  Best Local Similarity   100.0%;  Pred. No. 3.5e-280;
  Matches 1386;  Conservative    0;  Mismatches    0;  Indels    0;  Gaps    0;

Qy    166 AATGTTAAAGTAGAGACTCAGAGTGATGAAGAGAATGGGCGTGCCTGTGAAATGAATGGG 225
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db      1 AATGTTAAAGTAGAGACTCAGAGTGATGAAGAGAATGGGCGTGCCTGTGAAATGAATGGG 60

Qy    226 GAAGAATGTGCGGAGGATTTACGAATGCTTGATGCCTCGGGAGAGAAAATGAATGGCTCC 285
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     61 GAAGAATGTGCGGAGGATTTACGAATGCTTGATGCCTCGGGAGAGAAAATGAATGGCTCC 120

Qy    286 CACAGGGACCAAGGCAGCTCGGCTTTGTCGGGAGTTGGAGGCATTCGACTTCCTAACGGA 345
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    121 CACAGGGACCAAGGCAGCTCGGCTTTGTCGGGAGTTGGAGGCATTCGACTTCCTAACGGA 180

Qy    346 AAACTAAAGTGTGATATCTGTGGGATCATTTGCATCGGGCCCAATGTGCTCATGGTTCAC 405
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    181 AAACTAAAGTGTGATATCTGTGGGATCATTTGCATCGGGCCCAATGTGCTCATGGTTCAC 240

Qy    406 AAAAGAAGCCACACTGGAGAACGGCCCTTCCAGTGCAATCAGTGCGGGGCCTCATTCACC 465
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    241 AAAAGAAGCCACACTGGAGAACGGCCCTTCCAGTGCAATCAGTGCGGGGCCTCATTCACC 300

Qy    466 CAGAAGGGCAACCTGCTCCGGCACATCAAGCTGCATTCCGGGGAGAAGCCCTTCAAATGC 525
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    301 CAGAAGGGCAACCTGCTCCGGCACATCAAGCTGCATTCCGGGGAGAAGCCCTTCAAATGC 360

Qy    526 CACCTCTGCAACTACGCCTGCCGCCGGAGGGACGCCCTCACTGGCCACCTGAGGACGCAC 585
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    361 CACCTCTGCAACTACGCCTGCCGCCGGAGGGACGCCCTCACTGGCCACCTGAGGACGCAC 420

Qy    586 TCCGTTGGTAAACCTCACAAATGTGGATATTGTGGCCGAAGCTATAAACAGCGAACGTCT 645
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    421 TCCGTTGGTAAACCTCACAAATGTGGATATTGTGGCCGAAGCTATAAACAGCGAACGTCT 480

Qy    646 TTAGAGGAACATAAAGAGCGCTGCCACAACTACTTGGAAAGCATGGGCCTTCCGGGCACA 705
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    481 TTAGAGGAACATAAAGAGCGCTGCCACAACTACTTGGAAAGCATGGGCCTTCCGGGCACA 540

Qy    706 CTGTACCCAGTCATTAAAGAAGAAACTAAGCACAGTGAAATGGCAGAAGACCTGTGCAAG 765
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    541 CTGTACCCAGTCATTAAAGAAGAAACTAAGCACAGTGAAATGGCAGAAGACCTGTGCAAG 600

Qy    766 ATAGGATCAGAGAGATCTCTCGTGCTGGACAGACTAGCAAGTAATGTCGCCAAACGTAAG 825
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    601 ATAGGATCAGAGAGATCTCTCGTGCTGGACAGACTAGCAAGTAATGTCGCCAAACGTAAG 660

Qy    826 AGCTCTATGCCTCAGAAATTTCTTGGGGACAAGGGCCTGTCCGACACGCCCTACGACAGT 885
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    661 AGCTCTATGCCTCAGAAATTTCTTGGGGACAAGGGCCTGTCCGACACGCCCTACGACAGT 720

Qy    886 GCCACGTACGAGAAGGAGAACGAAATGATGAAGTCCCACGTGATGGACCAAGCCATCAAC 945
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    721 GCCACGTACGAGAAGGAGAACGAAATGATGAAGTCCCACGTGATGGACCAAGCCATCAAC 780

Qy    946 AACGCCATCAACTACCTGGGGGCCGAGTCCCTGCGCCCGCTGGTGCAGACGCCCCCGGGC 1005
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    781 AACGCCATCAACTACCTGGGGGCCGAGTCCCTGCGCCCGCTGGTGCAGACGCCCCCGGGC 840

Qy   1006 GGTTCCGAGGTGGTCCCGGTCATCAGCCCGATGTACCAGCTGCACAGGCGCTCGGAGGGC 1065
          ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    841 GGTTCCGAGGTGGTCCCGGTCATCAGCCCGATGTACCAGCTGCACAGGCGCTCGGAGGGC 900

Qy   1066 ACCCCGCGCTCCAACCACTCGGCCCAGGACAGCGCCGTGGAGTACCTGCTGCTGCTCTCC 1125
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                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     901 ACCCCGCGCTCCAACCACTCGGCCCAGGACAGCGCCGTGGAGTACCTGCTGCTGCTCTCC 960

Qy    1126 AAGGCCAAGTTGGTGCCCTCGGAGCGCGAGGCGTCCCCGAGCAACAGCTGCCAAGACTCC 1185
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db     961 AAGGCCAAGTTGGTGCCCTCGGAGCGCGAGGCGTCCCCGAGCAACAGCTGCCAAGACTCC 1020

Qy    1186 ACGGACACCGAGAGCAACAACGAGGAGCAGCGCAGCGGTCTTATCTACCTGACCAACCAC 1245
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1021 ACGGACACCGAGAGCAACAACGAGGAGCAGCGCAGCGGTCTTATCTACCTGACCAACCAC 1080

Qy    1246 ATCGCCCGACGCGCGCAACGCGTGTCGCTCAAGGAGGAGCACCGCGCCTACGACCTGCTG 1305
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Qy    1426 ATCCACATGGGCTGCCACGGCTTCCGTGATCCTTTTGAGTGCAACATGTGCGGCTACCAC 1485
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Db    1261 ATCCACATGGGCTGCCACGGCTTCCGTGATCCTTTTGAGTGCAACATGTGCGGCTACCAC 1320

Qy    1486 AGCCAGGACCGGTACGAGTTCTCGTCGCACATAACGCGAGGGGAGCACCGCTTCCACATG 1545
                ||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||||
Db    1321 AGCCAGGACCGGTACGAGTTCTCGTCGCACATAACGCGAGGGGAGCACCGCTTCCACATG 1380

Qy    1546 AGCTAA 1551
                ||||||
Db    1381 AGCTAA 1386
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Job time: 6786 sec